SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22299	52667	A	22427	9	1535	EAQQLVSERPAENRSWKCLLA ARGEERGASIMAEQOVENDL. DVDEEEEPQAPQESTPAPPKKD IKGSYWSIHSSGFRDF/LCLKPE LLRAIGGLVAFEHPSEVQIECCI PQGHPGAWDVLLPGQSSGMCK TAVFRCWPPLQNEPVNGQVT. VLMMCHTRELAFQISKEYERS KYMPSVKVSVFFGGLSIKKDVE EVMIKKNCPHVVYDFFGRILAL VRNRSFSLKNVKHFVLDECDK MLEQLDMRRDVQEIFRLTHIEK QDPMEVFVDDETKLTLHGLQQ YYVKLKDSEKTRKVFDLWDV LEFNQVDLRQGQCKRCMALA QLLIVEQNIFPAGWPHIRGNGFQ VATQILIFGIRGMDIEAKSINIVEN VAMPFEGLGFYLHRKVARGE FGTKRP*PFTFVSDGNDAIKILN
	50.00		22.422		200	DV\QDRF*S*MWAEL\PEEIDIST YIEQSRLNHQLARAGPPGAAPH AASPPLSRRHC
22300 22301	52668 52669	A	22428	3	290 690	MNQTSEKEIISKEQGNVKAAGA SDVVLYKIDVPANRYDLLCLEG LVRGLQVFKERNFFLTVLEVEK SKTKVEQIQCLCSQEDIADKLG VDISATKAVHISNPKTAEFQV KTILLPGLLKTHANRKMPLA KTILLPGLKTHANRKMPLA KLEEISDIVIKDSNTDVGAKNVR HLCAVYYNKNPGFEIHGLLDRI WQLLDVPPGEDKGGYVIQSVG KLGVLHPDVITKFELTMPCSSL EINIGPFL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22302	52670	A	22430	2	1834	GTRPTVSVKRDLLFQALGRTYT
22302	52070	ļ.,	22.50	ľ	1.05	DEEFDE\LCFEFGLELDEITSEKE
		i				IISKEOGNVKAAGASDVVLYKI
1		1		l		DVLATRYDLLCLEGLVRGLOV
		l		ĺ		FKERIKAPVYKRVMPDGKIOKL
		l		ŀ		IITEETAKIRPFAGAAVLRNIKFT
		1		İ		KD\RYDSFIELQEKLHQNICRKR
		l		ŀ		ALVAIGTHDLDTLSGPFTYTAK
		1		ł		RPSDIKFKPLNKTKEYTACELM
		1		ł		NIYKTONHLKHYLHIIENKPLYP
		1		l		VIYDSNG\VVLSMPPIINGDHSRI
		1		l		TVNTRNIFIECTGTDFTKAKIVL
l		l				DIIVTMFSEYCENQFTVEAAEV
		l				VFPNGKSHTFPELAYRKEMVR
ŀ				l		ADLINKKVGIRETPENLAKLLT
ŀ				i		RMYLKSEVIGDGNOIEIEIPPTR
				l		ADIIHACDIV\EDAAIAYG\YNNI
		l		l		\QMHSPENFTP*ANQFPLNK\LT
		ı				ELLRHDMA\AAGFH*STLPLAL
		l		İ		CSQGRYWLDKLGVGYLWQQR
		l				AVHISN\PKTAEFQVARTTLLPG
		l		l		PP*RP*AANRKV\PLPL\KLFEIS\
		l		i		DIVIK\DSNT\DVGAKKTTROSL
		ı				VAGLLPTKNPG\FEINSWGCWD
	ĺ	l				RIMAVCSDV\PPGGRQGGDM*F
		1				KASEGPAFFPGAMCRDSFARGQ
		ł				SL\GKLG\VLHPDVIHQIWS*PLP
		ł				CSSLEI\NIGPFL
22303	52671	A	22431	I	304	CSSLEIMIGFFE
22304	52672	A	22432	1	642	
22305	52673	В	22433	1	423	
22306	52674	A	22434	i	525	LHEHPVTESPMNCSVTLOLSLY
22300	32074	1	22737	ľ	323	MKSVA/GIGOTGKIEPVVLPLL
		1				WFAESGAMEGETLHTFYTOLV
		1				LMPKVMHYAQYVLLALGCVL
		ı		i		LLVPVICOIRSO/DRCWPEGSPG
		ı				*QPFAFPPA*TDIILWTEPATLM
		ı				AEVSACCAPELPPPPHRVPQIIG
		1				AQAEVNEPGGPAFRALLDPGA
22307	52675	A	22435	3	483	VRGSGNSHGEVSKGRMCLAGA
22301	12073	l^	12733	ľ	100	LPLPGACAALVLLLVILPEPLRC
		1		i		PLHTLYGGRDPVDSIRLWRQA
		l				GEFRHKSNITFNNNDTVSFLEY
	l	1	l	l		RTF\QFQPSKSHGSESDYIVMPN
		1		l		ILVLNKPRSTWISVCFVHGCIPA
	l	1	1	1		SRTGPRTESMHSKSLWKAWLA
		1	l	l		APGLPLL
		ட	L	L		AFGEFEE

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last aminu acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22308	52676	Α	22436	1	1623	RRRPRLLPGAEPCEPRVGPRRA
		l				DMGCSAKARWAAGALGVAGL
						LCAVLGAVMIVMVPSLIKQQV
	ŀ					LKNVRIDPSSLSFNMWKEIPIPF
						YLSVYFFDVMNPSEILKGEKPQ
						VRERGPYVYK/ESSRHKSNITFN
		1				NNDTVSFLEYRTFQFQPSKSHG
		l				SESDYIVMPNILVLGV\AVMME
		l				NKPMTLKLIMTL\AFTTLG\ERA
		ł				FHGTALWGEV\MWGYKDPLVN
		l				LINKYFPGMFPFKGGFGVFAEL
		1				NNSDSGLFTVFTGVQNISRIHLV
		l				DKWNGLSKVDFWHSDQCNMI
		l				NGTSGOMWPPFMTPESSLEFYS
		ı			l	PEACRSMKLMYK\ESGVFEGIP
		l				TYRFVAPKTLFANGSIYPPNEGF
		l				CPCLESGIQNVSTCRFSAPLFLS
	1	l				HPHFLNADPVLAEAVTGLHPN
	ĺ	1				QEAHSLFLDIHPVTIIPMNCSVK
		l				LQLSLYMKSVAGIGQTGKIEPV
	1	l		ŀ		VLPLLWFAESGAMEGETLHTF\
		1				YTQLVL\MPKVMH\YAQYVPS
		1		l	ł	WALG\CVLLLVPV/LSCQIRSQE
		l			1	KCYLFWSSSKKGSKDKEAIQA
		l			1	GSE*SLMTSAPKGS\VLQEAKL
22309	52677	Α	22437	198	1062	STCVEMPRNLLYSLLSSHLSPHF
	l	l		ľ		STSVTSAKVAVNGVQLHYQ\HF
		l				GEGDHAV\LLLPGMLGSGETDF
		1		l		GPQLKNLNKKLFTVVAWDP\R
		1		İ		GYGHSRPPDRDFPADFFERDAK
		1		İ		DAVDLMKALKFKKVSLLGWS
		1				DGGITALIAAAKYPSYIHKM\VI
		1				WG\ANAYVT\DERPAG*YEGHP
		ı	į			RMVSQMGVEGKKESL*EALYG
	1			l	1	YDYFARTCEKWVDGIR\QFKHL
		1		I		PDGNICRHLLP\RVQ\CPPPLIVH
				1		G\EKDPLVPRFHADFIHKHVKG
		l		l		SRLHLMPEGKHNLHLRFADEF
ĺ		1				NKLA\EDFLQ
22310	52678	A	22438	223	393	AOASPPLLPP**ERSNVPCPIS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22311	52679	Α	22439	524	1516	FILEDITGYVMAPIDGLLCREPR
		1		l	\	RGLYLEGTKPMTLFQIQFLNSV
		1				GDLLDLIPSLSPTKNGSLKVFKR
		ı				WDMGHCSALIKVLPGFENILFA
		l l				HSSWYTYAAMLRIYKHWDFNII
		l				DKDTSSSRLSFSSYPGFLESLDD
						FYILSSGLILLQTTNSVFNKTLL
		i i				KQVIPETLLSWQRVRVANMMA
						DSGKRWADIFSKYNSGTYNNQ
		1				YMVLGPEESKAEPQSLTKGT/P
		l				VTLWEQIPTY/VRNISGTNLMFL
		l		1		RGKGFWA\SYNVPFHEKIYNWS
İ		1				GLSHC*FRKLGLGPTPYDFRPH
ļ		1				EPKIFRRDPR/DK*LIRHPMKYI
						MRYN\NYKE\DPYSRGDPPVNT
22312	52680	Α	22440	1	306	
22313	52681	Α	22441	698	1613	FOFFOIFPEYKNNDFYVTGESY
						AGKYVPAIAHLIHSLNPVREVKI
		1				NLNGIAIGDGYSDPESIIGGYAE
l						FLYQIGLLDEKOKK\YFQKOCH
				1		ECIEHIRKONWFEAFEIL\DKLL\
		ı		1		DGDLTSDPFLTSONVTGCS\NY
Į.		ŀ		1		YNFLRCTEPEDQLYYVKFLSLP
						EVRPAHPTLGN\OTFN\DGPIV*
		l				KVPCGEDTSTSPVKPW\LTEI\M
						NNYRVLIY\NG\QLGHPSVAAAP
		ŀ				DOSAPLMG\MDWKGSP\EYKK
						AEKK\VWKIFKS\DSEVA\GYI\R
		1		i		OVGDF\HQVIIRGGGHILPYDOP
		l		İ		LRAFDMINRFIYGKGWDPYVG
22314	52682	Α	22442	1	258	
22315	52683	Α	22443	366	848	
22316	52684	A	22444	1	585	ECLCVSTAVRIRHELKTMLTEL
		1	l			LRGGSFEFKDMRVPSALVTLH
	1	1	İ	i		MLLCSIPLSGRLDSDEQKIQNDI
		1		l		IDILLTFTQGVNEKLTISEETLA
			l	l		NNTWSLMLKEVLSSILKVPEGF
			l	1	1	FSGLILLSELLPLPLPMQTTQVS
		1	l	l		LPYNMHLINDCSNTF*KASDSV
			l	1		KKQKPSSPFLPKRHKITG*GGD
l	1	1		1		KTETSWSLRKYGGKNFK

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ĺ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22317	52685	A	22445	625	2700	YIRGHSTPRKLWSMHLHVQAK
			ŀ			LLQEIVRSFSGTTCQPIQHMLRR
			ŀ		ľ	ICVQLCDLASPTALLIMRTVLD
			ŀ			LIVEDLQSTSEDKEKQYTSQTT
						RLLALLDALASHKACKLAILHL
			İ	ŀ	ļ	INGTIKGDERYAEIFQDLLALVR
				İ		SPGDSVIRQQCVEYVTSILQSLC
				1		DQDIALILPSSSEGSISELEQLSN
			ŀ		}	SLPNKELMTSICDCLLATLANS
	l					ESSYNCLLTCVRTMMFLAEHD
			ŀ		1	YGLFHLKSSLRKNSSALHKLLK
				ŀ	ŀ	RVVSTFSKDTGELASSFLEFMR
				ŀ	ĺ	QILNSDTIG\CCG\DDNG\LMGSR
			ļ.		ĺ	RGAHTSRTMSINAAELNQLLQS
			l	1		TEESPEDLFLELEKLDLEHSKD
				1		DDNLYSLLD**RGLKQMLESSG
				l		DPLPLSDQDVEPVLSAPESLQN
				İ		LFNNRTAYVLADVM\DDQLKS
				l		MWFTPFQAE\EIGTDLDLVKVD
				l		LIELSEKCCSDFDLHSE\LERSFL
				1		SEPSSPGRTKTTKGFKLGKHKH
				1	!	ETFITSSGKSEYIEPAKKEFMLC
				1	i	PPPRGRGRGGFGOGIRPHDIFR\
				1		ORKONTSRPP\SMHVDDF\VAA
			İ	İ		E\SKEVVPQDGI/PSHPNRPLKVS
1			1	1		ORISSRVGFSGNRGGRGAFHSO
				1		NRFFTPPASKGNYSRREGTRGS
1				1		SWSAONTPRGNYNESRGGOSN
1		ļ		1		FNRGPLPOL\RPLKFPTG*PAPK
1		ļ		1		SLGTRAS*MVRGG\LGTFPGLS
1	ŀ			1	ì	AKLAASGGSRGKFVSGG\SGRG
22318	52686	A	22446	2487	3161	TWWLHISHIPGGKKSYSPRLGR
22510	32000	^	22770	2407	5.01	GKRFATFVLLVGGELCLTSYAP
ł				1		PSIHPILRSEAECLNYSIFSKFLA
]				1	!	PAGEFALPRSFDNCPRVRLRAL
1				1		RLALLILLTFILCWTPYYLLGM
				1		WYWFSPH\MLTEVPPSLSHILFL
1				l	l	LGLLNAPLDPSPSYGAFTLGCR
1	l			l		RGHQELSIDSSKEGSGRMLQEEI
	l		l	l		
	l		1	l	I	HAFRQLEVQKTVTSRRAGETK
22216	50607	-	22447	222	227	GISITSI*S*QSM*EQNSKSLVP
22319	52687	A	22447	232	337	
22320	52688	A	22448	I	1449	
2232 I	52689	Α	22449	1	2004	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
22322		A	22450	3	1558	APARDRAGGGRRALPRIPAAS ARARGITRALSRAMACRQEPQ PQGPPPAAGAVASYDYLVIGGG SGGLASARRAAELGARAAVVE SKHLGGTCVNVGCVPKKVMW NTAVIISEFMHDHADYGFPSCE GKFNWRVIKEKRDAYVSRLNA IVQNNLTKSHIEIIRGHAFFYSD PKPTIEVSGKKYTAPHILIATOG WYPSTPH-FERSRGAS.I-YITSDG FFQLEEIVPAAASLVGARLPLSV EMAGILISALGSKTSMIRHDIX VLRSFDSMISTNCTEELENAGV EVLKFSQVKEVKKALISGLEAS WYTAVPGRIPVMPMIPDVDCL LWAIGGVPNTKDLSLNKLGIQT DDKGHIIVDEFQNTNVKGITAV BODVCGKALILTPVAIHAGRKLA HRLFEYKEDSKLDYNNIPTVVS SHPPIGTVGLTEDEAHKYGIEN VKTYSTSFTPMYHAVTKKKTK CVMKMVCANKEEKVVGIHM CULCOEMILOGFAVAVKMGAT
		L		-		KADFDNTVAIHPTSSEELVTLR
22323	52691	A	22451 22452	51	464	
22324 22325	52692 52693	В	22452	103	1184 148	
22323	52694	A	22454	1	3282	
22327	52695	A	22455	145	436	
22328	52696 52697	A	22456	3	3622	MPGYRAGPFEEAFDREDERQIE CQGRGVDRKITGEHITTFSGRSG PDHEITLIDSGGGFTRSRA ASEDRRKLPIIVEDEGGFTRSRA CSSPARGSRPPPSAIGCSPVAQA SDSAAGPARRTALQSLSSWLGY QIDRHSVPVYVFKSPLFSVIMAP KHKSSDAGDLDPRKRSKVLPL SEKVKVLDLIRLDKKSYAEVA KIYGKNESSIREIVKKEKEIRASF EQALHLWVEE VVELRCQDKEGDPAPEYTWFK DGIRLLENPRLGSQSTNSSYTM NTKTGTLQFNTVSKLDTGEYSC EARNSVGYPOVPGKRMQVDD
22330 22331	52698 52699	A A	22458 22459	I	20 <b>8</b> 321	LNISGIIAAVCGLGVCYAQRKG YFSKETSFQKSNSSSKATTMSE NDFKHTKSFII

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22332	52700	A	22460	2	1169	HTPKTEQTPIPGLEDPPLGSQLR
				l		RRPGEGETDIPSRAVPPLPPLPT
				1		LCSFPAPEVQGPPASCAPAAGT
				l		LDLLRAAGCRPGKMARRSRHR
				l		LLLLLRYLVVALGYHKAYGF
						SAPKDQQVVTAVEYQEAILAC
		ŀ		l		KTPKKTVSSRLEWKKLGRSVSF
				l		VYYQQTLQGDFKNRAEMIDFNI
		i				RIKNVTRSDAGKYRCEVSAPSE
1				l		QGQKPG\EDTVTLEVLGDVHVL
1						APAVPSCEVPSSALSGTVVELR
1				l		CQDKEGNPAPEYTWFKDGIRLL
		ĺ		l		ENPRLGSQSTNSSYTMNTKTGT
				l		LQFNTVSKLDTGEYSCEA\RNS
				l		VG*SEGPCETNASR*SQHKVGII
		l		I	1	GSRS*LWALVDFPFVGLVYSYG
		l		I	1	SEGKATFSKETSFQKSNSSSKAT
						TMSENDFKHTKSFII
22333	52701	В	22461	57	168	
22334	52702	Α	22462	1	682	
22335	52703	Α	22463	398	1465	GTFPPSGLITGKKRKALLGACD
		ĺ				CTQIVKPSGVHLKLVLRFSDFG
		l				KAMFKPMRQQRDEETPVDFFY
1						FIDFQRHNAEIAAFHLDRILDFR
				1	1	RVPPTVGRIVNVTKEILEVTKN
				l	1	EILQSVFFVSPASNVCFFAKCPY
i i				l		MCKTEYAVCGKPHLLEGSLSA
İ				1		FLPSLNLAPRLSVPNPWIRSYTL
				l		AGKEEWEVNPLYCDTVKQIYP
		l				YNNSQRLLNVIDMAIFDFLIGN
				l	-	MDRHHYEMFTKFGDDGFLIHL
				l		DNA\RGFGRHSH\DEIPILSLLSQ
				1		CGMIKKNTLLHLQ\LLAQADYR
		l		1		LS\DVMRESLLEDQL\SPVLTEP
						HLLAL\DRRLQTILRTV\EGCIV\
						AHG\QQSVH*FDGPSGNSSGPR
22336	52704	Α	22464	2	424	GRVGVSAVIVRAWITHKYSGR
İ		l		1		DWNVKWEENLLHAVAKNYTL
		l		1		LQTIPPFERPFKDHQVCLEWNM
1		l		1		GYIWNLRANRIPOCPLEN/DFRE
		l		i		DFHYNDTAGYFIIGGSRYVAGI
		l		I		EGFFGPLKYYRLRSLHPAQIFNP
				l		LLEKOLAEOIS
22337	52705	A	22465	23	323	
22338	52706	A	22466	3	287	LSVQMYAQAALDGDSQGFFNL
				ľ	Γ".	ALLIEEGT\YERCWSHSNEESFS
		ĺ		l		PCSLAWLYLHLRLLWGAILHSA
1		l		l		LIYFLGTFLLSILIAWTVQYFQS
1				l		VSGKDL
22339	52707	A	22467	1	348	.55.122
22340	52708	A	22468	1	738	
22340	122700	<u></u>	22400	L <sup>1</sup>	1,20	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22341	52709	A	22469	1	507	
22342	52710	Α	22470	187	3682	SILCLLSPCVVOFGKPVVSILSS
						RSRHSPCTKKGWEGMRKHLHT
				1		ROGHK*VHV\KFP\SIMGLOGM
						DYFHPDISIFCNLAVIVRAWITH
			į .			KYRGRDWNVKWEENLLHAVA
						K\NYTLLQTIPPFERPFKDHQVC
						LEWNMGYIWNLRANRIPOCPL
						ENDVVALLGFPYASSGENTGIV
				i		KKFPRFRNRELEATRRQRMDYP
						VFTVSLWLYLLHYCKANLCGIL
				1		YFVDSNEMYGTPSVFLTEEGYL
						HIQMHLVKGEDLA
22343	52711	A	22471	3	153	
22344	52712	Α	22472	1	408	TEPEFPGRRFRPRMAVMAP\RT
1			l			LVLLLSGALALTQTWAGSHSM
						RYFSTSVSRPGRGEPRFIAVGYV
1		1	l			DDTQFVRFDSDAASQRMEPRA
				1		PWIEQEGPEYWDEETGKVKAH
l			l			SQTDRENLRIALRYYNQSEAGS
						HTLFPR
22345	52713	Α	22473	]1	823	MAVVAPRTLLLLLSGALALTQT
			İ			WAGSHSMRYFSTSVSRPGSGEP
						RFSAVGYVDDTQFVRFDSDAA
l			İ			SQRMEPRAPWMEQEEPEYWDR
			l			QTEISKTNAQIDLESLRIALRYY
						NQSED/GPPPKTHMTHHPISDHE
				1		ATLRCWALSFYPAEITLTWQRD
				1		GEDQTQDTELVETRPAGDGTF
						QKWASVVVPSGQEQRYTCHVQ HEGLPKPLTLRWEPSSQPTIPIV
				1		GILAGLVLFGAVIAGAVVAAV
				1		MWRRKSSDRKGGSYSQAASSD
						SAOGSDVSLTACKV
L	l		L	l	L	3AQU3DV3LIACKV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequenec ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence	ı	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		i		sequence		
22346	52714	A	22474	1	1162	NSRVDDFVAVMAPRTL\LLLLS
22340	32714	<u> </u> ^	22474	ľ	1102	GALALTOTWAGSHSMRD\FFTS
1	ŀ					VSRPG\R*EPRFIAVGYVDDTOF
					1	VRF\DSDAASQRMEPRAPWIEO
					ł	EGPEYWDQETRNMKA\HSQTD
1						RENLRIALPLTYNOSEAGS\HTL
						QMMFGCERGGSDGALSFRGLP
					1	PVTAY\DGKGFTFALNE\DLRS
	ĺ					
						WT\AADMAAQITQRKWETPPM
				1		WRS/EQEEAYLEG\QCV\DGLRR
						YL\ENGKEDAAAQRTPPKTHMT
	i			1	l	HHPIS*P*RPL*RCW\AL\AFYPC
						GDHT*PWQRDG\EDQTQ\DTEL
						V\ETRPAGDGTF\HKWAAVVVP
			ĺ			SGEESEITPANVQQ*RVLPKPLT
						LKM/WEPSSPAHPSPIVGHHLLA
						LVL/LLGAFDHLKLWVAAVML
		l				KEGTAQVGKG*KIEKGGSYSSR
						LPSQ
22347	52715	Α	22475	9	316	IPAMGLETEKADVQLFMDDDS
1		l			\	YSHHSGLEYADPEKFADSDQD
		l				RDPHRLNSHLK\ILMPFVKTCL
		l				MVLPSVQTIWKSVTDVIIAPLCT
			1			SVGRCFSSVSLQLSQD
22348	52716	A	22476	3	606	RPRTKAAMGLETEKADVQLFM
		ł				DDDSYSHHSGLEYADPEKFADS
						DQDRDPHRLNSHLKLGFE\DVI
						AEPVTT\HSFDKVW\ICSHALFEI
						\SKYRM\YRSLT\VFLGHSPWAFI
				1		CGEFSFATL\SCLAHLGF*WPFV
						KTCLM/VFLPSVQTIWEEC*QM
	İ			1		V/ISLAPLC\TSVG\RCFLFLFSLA
	l					TEPRNGILGPOVWRLGYCNTSF
						VIIT
22349	52717	Α	22477	101	432	ATROQLKAKPENLLGSISRDSG
				1		RTOEGIPARVPYPPPP/PTREPPO
						ST*PGRNGLLGCLGGGAPRPG
						WLDSEAGGAPAFHCPG*CPRRH
						RRGGTRLKOLPPPPTTVHDOPL
22350	52718	A	22478	794	1209	VKVDEVTVKHLGCORCOWSRC
	[	ľ.	I	l	1	PPSCSGTCLCCGAEAPCAAOPA
		1	1	l		WPPPPAAPPPTPPAPPSD\NSSCT
			l			O*PTPRCGT*DKDKGIFTLPKGH
						P*V*SLEHGFLKLGSCP\PPPPHA
				l		LVRPPOVRFNTPHGAEPORDOL
		1	1	l		, , ,
		Ц	L	L	L	PQST

SEO ID	SEQ ID NO:	Mat	SEO ID NO:	Nucleotide	Nucleatide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
		<u> </u>				
22351	52719	A	22479	3	1041	GSNCPLRQADPGPPRHTLFMDF
		1				VAGAIGGVCGVAVGYPLDTVK
		1				VRIQTEPKYTGIWHCVRDTYHR
		l				ERVWGFYRGLSLPVCTVSLVSS
		1			1	VSFGTYRHCLAHICRLRYGNPD
				l		AKPTKADITLLGCGLRTRPRVP
		l	i			DVAH*GGQSPLADADTGFLSSS
1				l		GGFRLSALLAAPPMCPVPPACP
į.					1	EPKYRGPLHCLATVAQ\EEGLC
ŀ						GLYKGSLALVLRDGHSFATYFL
		1				SYAVLCEWLSPAGHFRPDVPG
		1		1		VLVAGGCAGVLAWAVATPM\D
						GIKSRLEANGQGQ\NGYRGLLH
					İ	CMVTSVREEGPRVLFKGLVLN
ļ		l				CCRAFPVNMVVFVAYEAVLRL
						RPGSAHHSRSPTPRGATHQQLV
22352	52720	Α	22480	38	453	
22353	52721	A	22481	287	413	DMDYRRGGWTVIHRKIGGRID
						FQRVR*DYLHGYADLLREFWL
22354	52722	Α	22482	1	393	MPKLLKWNYEGGACWGNLW
						KLPNVERLCFSEATVFRSAAQC
1					Į.	LVVGLGLLLVSRACSHEDAIEW
1						RGDSSVVNIVEDGSNAKDESKS
		1				NDTVCKEDCEESCDVKTKITRE
1			Ì		ĺ	EKHFMCIFHNVQSLGTVCGEK
						GIDVN*NYTRRKTFHVYLPQCS
						VTGNSLRGERH
22355	52723	A	22483	505	854	
22356	52724	A	22484	584	1009	FTQKDLFIPFEVMCDMDYRGG
1						GWTVIQKRIDGIIDFQRLWCDY
1					İ	LDGFGDLLGDAFL\FL*KEDNQ
					ŀ	NAMPFSTSDVDNDGCRPACLV
1						NGQSVKSCSHLHNKTGWWFN
						KCGLANLNGIHHFSGKLLATGI
						QWGTGNQKQLTWQE
22357	52725	A	22485	1	3123	MSHRARPCPAPFLFWMLGVKT
						EPAGQDRRKKARSRCCIFRVHA
		l	ĺ			SGLWGLRFPSVEWGKPSRDKA
						QLSESPPQSVYLFAETALSAIVP
						GDGVAGGGGTGDGGPGAPYPG
						AVADMEELTIWEQHTATLSKD
			l	l		PRRGFGIAISGGRDRPGGSMVV
		l		Į.		SDVVPGGPAEGRLQTGDHIVM
1	1	l	l	l		VNGVSMENATSAFAIQILKTCT
	1	l				KMANITVKRPRRIHLPATKASP
1	1		l	l		SSPGRQDSDEDDGPQRVEEVD
	1	ŀ	1			OGRGYDGDSSSGSGRSWD
22358	52726	Α	22486	191	267	
						·

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first coden for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		097.740,217	sequence	or pepulae sequence	decision, - possible nucleotide inscrition)
		$\perp$				
22359	52727	Α	22487	2	609	PPPRSSEDPAKMAEVEHEEGSG
						PFRKFTLPAAVDL\DQ\RLDISY
		1				DQLEQPMQAVPVARPAAGGLN
		1				PGACGRKQ\HSPA*KRLLQGPS
		l				KGGAAPCHKPEAGEDATLRD\
		1				MII\LPEMWGS\MVGVYH\GKTF
						\NQVEDQGVCGRPCRLGWAGV
		1				A*CRRDQLTPSFALGLPQPEMI
		1				GHYLGEFSITYKPVKHGRPGIG
						ATHSSRFIPLK
22360	52728	Α	22488	3	461	STISLYCGLKTCPKLTCAFPVSV
1		ŀ				PDSCCRVCRGDGELSWEHSDG
1			ŀ			DFFRQPATREASHSCHRSHYDP
		l			İ	PPSRQAGGLSRFPGARSHRGAL
i		l			1	MDSQQASGTIVRFVLNNKHKH
1						GQVCVSNGRTYSHGESWHPNL
		1				RAFGIVECVLCTPTPID\WPQGD
1		1				*GKCRHGCPTLHYVD*LPLEKQ
		l				VILATALTMILHQADRLEVCPA
1						FLGPEVTGELLWIPSKHQEPLC
		1				DLSSITNTSMDKCVFPMEGPIL
		l				MASPGTQTSGLLALWSVCYVL
		_				PPPSIVPQGD
22361	52729	A	22489	656	1495	TGDGELSWEHSDGDIFRQPANR
		1				EARHSYHRSHYDPPPSRQAGGL
					1	SRFPGARSHRGALMDSQQASG
						TIVQIVINNKHKHGQVCVSNGK
İ		1				TYSHGESWHPNLRAFGIVECVL
İ		1				CTCNVTKQECKKIHCPNRYPCK
1		1				YPQKIDGKC*QNSTCFLIPEELP
		1		i		GQSFDNKGYFCGEETMPVYES
		l				VFMEDGETTRKIALETERPPQVI
		l		1		CSTFCLA\QGILQHFHIEKISKR
		l		I		MFEELPHGHFAFVCLLGQWKIF
		l		l		TEGEAQISQMCSSRVCRTELED
000.00	50000	١.	22.400		400	LVKVLYLERSEKGHC
22362	52730 52731	A_	22490 22491	1	477 294	
22364	52732	A	22491	56	4042	VFFTSMGEHLQKLIPASPACLA*
22304	32/32	l <sup>A</sup>	22492	36	4042	
				1		IPQACQP*PPTPPAWQCRNLR/L AASRHOEPRGEVGGPLGHGAE
		1		1		RILAVPVRTDAQGRLVSHVVSA
	1	1	l	I		ATSRAGVRARRAAPVRTPSFPG
		1		I		GNEEEPGSHLFYNVTVFGRDLH
	i	l		1		
	l	I	l	l		LRLRPNARLVAPGATMEWQGE
	l	1	l	l		KGTTRVEPLLGSCLYVGDVAG
	l	1	l	i		LAEASSVALSNCDGLAGLIRME
	l	1	l	I		EEEFFIEPLEKGLAAQEAEQGR
	l		l	1		VHVVYRRPPTSPPLGGPQALDT
	L	Ц.				GASLDSLDSLSRAL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequenee	of peptide sequence	deletion, \=possible nucleotide insertion)
22365	52733	Α	22493	173	1148	QALGLPSLLAVLALSTTDRASF
		1		\		SAVISRSPORTVSSTRKEORPLV
		l				FAFSALRLGTNACKMDGFCSTF
		l	İ			LGAPGIRLKLLTPAGTHKSERD
		l				RTATPTRHHEKRLFCSSLSIGRQ
		1				AEEAWRADHPLPESRLFGLVPL
		1		i		GGAAWCRLAGSFCVCSLRLPRP
		1		l		IELVRAASGALLGAVRRRNHGP
	1	1				PHCFSTATLLCKAYRGGHLTIR
		1				LALG\GAPNRPFYRIVAAQNKC
		1		ŀ		PRDGPFP*SKLGPPYDP/LLPPNS
	1		ŀ			HGEKTPCPLN\LDRI\RHWIG\CG
		l				APPLLSPMEKLLGLAGFFP\LHP\
						MMITNAERLRRKRAREVLLAS
			ŀ			QKTDAEATDTEATET
22366	52734	A	22494	2	4985	AGATTALELPGQPSVTGVPELP
						GLPSATRALELSGQPVATGALE
			ŀ			LPGPLMAAGALEFSGQSGAAG
						ALELLGQPLATGVLELPGQPGA
						PELPGQPVATVALEISVQSVVT
					İ	TSELSTMTVSQSLEVPSTTALES
						YN/DERSMMSSYERSMMSYER
			ľ			SMMSPMAERSMMSAYERSMM
						SAYERSMMSPMAERSMMSAYE
						RSMMSAYERSMMSPMADRSM
						MSMGADRSMMSSYSAADRSM
						MSSYSAADRSMMSSYTADRSM
22367	52735	A	22495	18	174	ALIKKHFIF*IAAAKPTPPKSQV
						TLTKEFPVSSGSQHRKKEADSV
				1		YGEWVL
22368	52736	Α	22496	3	595	IKKHFIF*IAAAKPTPPKSQVTLT
1		1				KEFPVS\SGPQHRKKEADSVYG
		l				EWVPVEKNGEENKDDDNVFSS
						NLPSEVSRRNIMYFSFFIPESAH
						FIVMLCLIWDSVSLFLRAGVKR
						QGRV*DGQM\KQPAASH\LTVT
						RM/CNSLCG\TKATKWKRLRFA
						ENSVI\TSLPNIGPSLHLWEGSP
						WYSIPFSFPSLLLLKVGEVEFKR

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown,  *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	l district			sequence		
22369	52737	A	22497	3	5618	SSELPGPSVTPVPQLSQELPGLP
						APSMGLEPPQEVPEPPVMAQEL
				l		PGLPLVTAAVELPEQPAVTVA
l						MELTEQPVTTTELEQPVGMTTV
	100	ĺ				EHPGHPEVTTATGLLGQPEATM
ŀ						VLELPGQPVATTALELPGQPSV
						TGVPELPGLPSATRALELSGQP
					ľ	VATGALELPGPLMAAGALEFS
						GQSGAAGALELLGQPLATGVL
					ŀ	ELPGQPGAPELPGQPVATVALE
			ļ.		ŀ	ISVQSVVTTSELSTMTVSQSLEV
				l		PSTTALESYNTVAQ
22370	52738	Α	22498	1	4685	MAPESHILASNTMETHILASNT
					l	MDSQMLASNTMDSQMLASNT
				l	ŀ	MDSQMLASSTMDSQMLATSSM
						DSQMLASNTMDSQMLASNTM
					İ	DSQMLASSTMDSQMLATSSMD
						SOMLATSSMDSQMLATSTMDS
						OMLATSSMDSOMLATSSMDSO
					İ	MLATSSMDSOMLATSSMDSOM
					[	LATSTMDSQMLATSTMDSQML
						ATSSMDSOMLASGTMDSOMLA
1						SGTMDAQMLASGTMDAQMLA
					ŀ	SSTQDSAMLGSKSPDPYRLAQD
					ŀ	PYRL
22371	52739	Α	22499	291	7655	KRAPEKVDAYLYKOILLMKLP
						TKKSKKHKKHKNKKKKKKKE
						KEKKYKROPEESESKTKSHDD
						GNIDLESDSFLKFDSEPSAVALE
l						LPTRAFGPSETNESPAVVLEPPV
l			1			VSMEVSEPHILETLKPATKTAE
1			l			LSVVSTSVISEQSEQSVAVMPEP
l			1	l		SMTKILDSFAAAPVPTTTLVLK
				l		SSEPVVTMSVEYOMKSVLKSV
l	l	l	1	l		ESTSPEPSKIMLVEPPVAKVLEP
				l		SETLVVSSETPTEVYPEPSTSTT
				l		
				L		MDFPESSAIEALR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
00000		<del> </del>		0.0	laree	COVERNI DEL CAMBRELLA COMO
22372	52740	A	22500	90	2166	SQHFGRLRRVDYLRTSKMQTSS
						SRSVHLSEWQKNYFAITSGICT
						GPKADAYRAQILRIQYAWANS
						EISQVCATKLFKKYAEKYSAIID
						SDNVESGLNNYAENILTLAGSQ
						QTDSDKWQSGLSINNVFKMSS
						VQKMMQAGKKFKDSLLEPALA
						SVVIHKEATVFDLPKFSVCGSS
1						QESDSLPNSAHDRDRTQDFPES
						NRLKLLQNAQPPMVTNTARTC
						PTFSAPVGESATAKFHVTPLFG
						NVKKENHSSAKENIGLNVFLSN
						QSCFPAACENPQRKSFYGSGTI
			ŀ			DALSNPILNKACSKTEDNGPKE
					i	DSSLPTFKTAKEQLWVDQQKK
						YHQPQRASGSSYGGVKKSLGA
1		1				SRSRGILGKFVPPIPKQDGGEQN
						GGMQCKPYGAGPTEPAHPVDE
						RLKNLEPKMIELIMNEIMDHGP
						PVNWEDIAGVEFAKATIKEIVV
1						WPMLRPDIFTGLRGPPKGILLFG
			ŀ			PPGTGKTLIGKCIASQSGATFFSI
						SASSLTSKWVGEGEKMVRALF
					1	AVARCQQPAVIFIDEIDSLLSQR
1	1					GDGEHESSRRIKTEFLVQLDGA
1						TTSSEDRILVVGATNRPQEIDEA
1						ARRRLVKRLYIPLPEASARKQI
1						VINLMSKEQCCLSEEEIEQIVQQ
1		1				SDAFSGADMTQLCREASLGPIS
1		l				AV/SQTADIATITPDQVRPIAYID
1		l				FENAFITVRPIVSPKDL*LYENW
22373	52741	Α	22501	3	523	FFFFFPAFKMSKRG\RGWVPSG
i	1	Į.				CRKFPDFPLGLP\VGTL*SNWCL
1		l	1			TNT\GAQKTLYI\ISREGGIKGTG
1		l	1			LTRLPRLPGVG\DMVMAHSSRK
1		l	l			ANPELQKKGYIPA\VVIRQRKVI
1		ĺ	I			TVGKRWRCFFIFED\NAGVIGD\
		l				NKG\EMKGFAITG\PRAKECAD
1		l	1			LWPRDCIQMLGSIAWFSQYIL
L			L	L	l	E W I KDCIQMEOSIA WISQIIE

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino seid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22374	52742	A	22502	13	946	GPSASVSFWLWSQQHSPRIRHP
22374	32/42	^	22302	13	340	RPGERRRLLHGIPHPTAEKOPA
		l				PLRPERRGGRRRRPAGRRGCA*
		l				/GGARPCSSRSS*PASTPAPRAA
		l				PPCRSPCSARPRTGPRTGAGRPR
		l				AP/HDGPPCATPDPRDOAS\PLPP
		l				RGPRPRVAPGSP\PEGHPRELPS
		1				EGSPGPRAPGORHVRPPRAVLR
		1				PRGLRGGGCPGPAHPAPRLQRR
						RAPGEAAAPGPRWPGRPGRGR
1		l				AOALVPAARSRCFWKLWEAVF
						CLPVAAAPTRRFGARORPOLW
		1				LHGHKGGOVGEVSSHSAGEEH
		1				LVPHAAPKRAREPHGRSGSPGG
		1				IASSWODGS
22375	52743	A	22503	34	823	IA33 WQDGS
22376	52744	A	22504	424	2339	LLRGTLQSYAWTSVIPDIATELR
22370	32/44	n	22304	1424	2339	DAPEKTLACMGLAIHQVTICLM
						HTFV*TFLMFAHTCTPCLTMVN
		l				VPHIHAR*GI*CIKVLLRMGH*R
					1	PFKNENAHRGTVVRVSNIKPLC
		1				TKMAFLCAACGEIQSFPLPDGK
		1				YSLPTKVIRSLTASLFILVKKAN
		1				QNTENISRQSIK*AIRLLHKRQA
						LDVTLLRKSIILLHDLVDSCVPG
		1				DTVTITGIVKVSNAEEGRVOLF
		1				SLFVSQ**FINVHLF*DAPPHILV
		1				VGDPGLGKSQMLQVDNQSFSV
		l				CSFACKMR**FTSEFSPLSKDSS
		1			1	SGDFALEAGALVLGDOGERPK
		1				GIISDSGTF*KAFSFSVNFLSISL
		ı				AKAGVVCSLPARTSIIAAANPV
		1				GGHYNKAKTVSENLK*VSSNIY
		1				TFNVLNHFNSIRKRNSSHLLSEH
		1				VIAIRAGKQRTISSATVARMNS
						ODSNTSVLEVVSEKPLSERLKV
						*MFLLLIQFGSV*MSKFSVWLIV
1	1	1			1	DVYPRLSTEAARVLQDFYLELR
						KQSQRLNSSPITTRQLESLIRLTE
		1	1			VCFFLWSCFFWLKGEGCALTSL
		1				LKYR*DKNFKLF*MKYOLSSHL
			1			KLGNRPNRSTAKRFISALNNVA
		1				ERTYNNIFQFHQLRQIAKELNIQ
		l				VADFENFIGSLND\QGYLLKKG
		1		1		PKVY\QLQTM
22377	52745	A	22505	73	214	I K T I V L V I W
22377	52746	A	22506	1	503	
223/6	32/40	ΙΛ.	122300	I <sup>1</sup>	303	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
				<u> </u>		
22379	52747	Α	22507	3	1000	ESKLEGKAAQDIKDEEPGDLGR
				ľ		PKPECEGYDPNALYCICRQPHN
						NRFMICCDRCEEWFHGDCVGIS
1			•			EARGRLLERNGEDYICPNCTIL
						QVQDETHSETADQQEAKWRPG
	ŀ					DADGTDCTSIGTIEQKSSEDQGI
						KGRIEKAANPSGKKKLKIFQPG
	İ					PGPVPTQLPVLWQVLEIAVSRSI
						SAFTLLHCISCKVIEAPGASKCI
	ŀ					GPG\CCHVAQP\DSVYC\SNDCIL
						KHAA\ATMKFLS\$GKEQKPKPK
1						EKMKMKPEKPSLPKCGAQAGI
1	ł					KISSVHKRPAPEKKETTVKKAV
						VVPARSEALGKEAACESSTP\S
1						WAS\DHNYNA\VKPEKTAAPSP
						VTVV
22380	52748	Α	22508	I	2330	MAAAEGKERAGGRASGRGRSE
	1					PPRPGARASDAGPGGLREVEGT
						GEHSVVAAGTVPRVGVGAAAR
						GEERRAGLGECGPGLGGCVAG
	}					LISGKSVQAFSVGAGRWPLLRV
						FGGNVTFCSVVSWSLCAYTLPL
						GMDDKGDPSNEEAPKAIKPTSK
	ł .			l		EFRKTWGFRRTTIAKREGAGD
	ļ					AEADPLEPPPPQQQLGLSLRRS
	1					GROPKRTERVEOFLTIARRRGR
						RSMPVSLEDSGEPTSCPATDAE
1						TASEGSVESASETRSGPQSASTA
						VKERPASSEKVKGGDDHDDTS
		1				DSDSDGLTLKELQNRLRRKREQ
						EPTERPLKGIQSRLRKKRREEGP
						AETVGSEASDTVEGVLPSKQEP
	İ					ENDQGVVSQAGKDDRESKLEG
						KAAQDIKDEEPGD\LGRPKPEC
						EGYDPNALYCICROPHNNRFMI
						CCDRCEEWFHGDCVGISEARG
		1		1		RLLERNGEDYICPNCTILQVQD
						ETHSETADQQEA*MGDLGDA\D
1		1		l		GTDF\TSIGNIRSRSLSRRPKGLK
						GRIEKAANPSGKKELKIF\OPVL
1		İ				EIAVSRSISAFTLLHCISCKVIEA
						PGASKCIGPGCCHVAQPDSVYC
						SNDCILKHAAATMKFLSSGKEO
						KPKPKEKMKMKPEKPSLPKCG
		1				AQPVQWLLLLSSWQLGTSSGL
		l				OVTLCOPPFTEHKWLKFSVLHG
						ETAGPPRAHRIRKLNKYWGAC
l						HHCMPLRPLPCLPGWPRSSPSL
	l			1		HILLWIFERPERCERO WERSSPSE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ļ				sequence	,	
22381	52749	A	22509	149	2001	I IQPPDPRNMTLAAYKEKMKELF
22301	32143	l^	22303	177	2001	LVSLFCSCFLADPLNKSSYKYE
						AGVGDSVGGRMKASGKTVLT
		1				GRERRAQADTV\DLNWCVISD
				1		
				1		MEVIELNKCTSGQSFEVILKPPS
						FDGVPEFNASLPRRRDPSLEEIQ
						KKLEAAEERRKRGPDLVIYEIH
						SPASLPFHSSDASVLGDIPPTTY
						SNISNSNQELLLMGGSPAPEQP
						HYEANILERSETTSYDNGSDAA
						QSFKVDGRCVFYINIPIPSNEGK
l					1	VLWKAPEIATAKEPTWSPLGST
						GPRPCQILVVKSPVVHGKGPVL
						SGCGLDLLEQPTFALCFLVAGA
				l		VTITFLSCSGEEAREESAPKAMS
ļ						ERQARRGGAEKQGAEGRGLQA
						VTTPELPYRMGLGAAFLLKERS
ļ		1				GVLPGSFICEDFTAALKQDEEIG
						TMKGKIREAYFKSFVPTPVAFS
ĺ				1		GGVMLPTGGTLEMCGNNFGGH
	1			1		KEQAGTVLPAFAIALRVTPSAS
l		1				LVLRPLNLDQAMLQGSLGLQL
					1	ADGLFWDLASIMDWTASRIQPS
		1				VSSSMKICQAFPGRPLAILLAQC
						AYVYGTWMKLEAIILTKLTQE
						QKTKHHLFSLPCCMPADISYRQ
						RCLGEPPHCTVWNRHLESLVM
		_				MSNSTTSPCSELLIFEVPPVAPE
22382	52750	Α	22510	373	990	IQPPDPRNMTLAAYKEKMKELP
						LVSLFCSCFLADPLNKSSYKYE
						ADTVDLNWCVIFRHWKVIE\LN
				İ		KCT/SGANSFEVILKPPSFDGVP
						EFNASLPRRARPIPWEEIPVRKL
						GRRLRERREVTREADP*KQLAE
ŀ		1		1		KR\EHEERGDSKRAFRETTTFIK
						MAK/EKNLAQKMESNRENREA
		1		1		HLAAMLERLQEK\DKHAEEVR
		L_				KNKELKEEASR
22383	52751	A	22511	524	1107	LCSPKVLGLNASCQKNFLKQK
		1	1			KSIKTLSTSQ/YSVSGYSLARSE
		1			1	NAHPEFRMAMKDGELVIWDSV
			1		1	HPCYTVFHEQTETFSSLWSEYH
						DDFRQFLHIYSQDVACYGENLA
						YFPKGFIENMFFVSANPWVSFT
		1		I		SFDLNVANMDNFFAPVFTMGK
		1	1		1	YYTQGDKVLMPLAIQVHHAVC
						DGFHVGRMLNELQQYCDEWQ
22384	52752	Α	22512	1	1413	
22385	52753	В	22513	1	2787	
22386	52754	Α	22514	1	1638	

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	coden for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22387	52755	A	22515	71	2421	YTVCYFLQAAEEDETQNQHDD
						SSSDSGTPSGPDFNYILNMSLW
					İ	SLTKEKVEELIKQRDAKGRE\V
					ľ	NDLKRK\SPSDLW\KEDLAGIC
						WRDWDKV/ENSPRNEEDVLAG
						MSGKSNLKGKVGKP\KVKKLO
						LEETMPSPYGRRIIPEITAMKAD
						ASKKLLKKK/KGVDLDTAAVK\
						VEFDEEFSGAP\VEGAGEEGIDS
						HQFPINK\GPKP\RREKKEPGS\R
						VRKTPTSSGKPSAKKVKKRNP
		1				WSDDESKSESDLEETEPVVIPR
	l	1				DSLLRRAADIHRLGGEGFDAGL
	1	1				TVFAPLFASPFPPRGAPYLKAA
		1				COTPRSGIGATAISGEALKGAV
		ł				SSGVDPRWASHLRVKLLEKSFS
		1				LQEIQSTEKVLSQQAKTLSLLG
		1				EPKLSKOLTGTCIYRKTNLALC
		l				RGPAAAKSLFOGAVRNDRNKK
		l	l			KKETSKQECTESYEMTAELDDL
		l	l			TEKIRKAHOETFPSLCOLGKYT
		l		l		TSLOKGMORCOILDLSSVRPIV
		ı				DVRPPELQDSKLVLVQAAKCA
		1		l		PLADHCSKPVLLKEGPEDOKD
		l		ĺ		QHDVGDCWNPECQTLFDQNN
		l	ŀ	1		AAKKEESETANKNDSSKKLSVE
1		1		1		RVYHIKTOLEHILLRPDTYIGSV
1		l		1		EPLSKVAEEGPAAVGGWKGWF
		l				GTESRKDPKPTGTEEVERGVW
1		l		1		LSGREGRLLSPSVVTTMVLAEN
		ı		1		SAIRFVGGKPSSDTVPKPKRAP
		l				KOKKVVEAVNSDSDSEFGIPKK
22388	52756	A	22516	1424	4860	GRDRYGVFPLRGKILNVREASH
22300	32/36	Ι^	22310	1424	4800	KQIMENAEINNIIKIVGLQYKKS
		l		1		YDDAESLKTLRYGKIMIMTDO
1		l		1		DODGSHIKGLLINFIHHNWPSLL
		l	1	l		KHGFLEEFITPIVKASKNKQELS
		1	İ	1		
		I	1	I		FYSIPEFDEWKKHIENQKAWKI KYYKGLG\TSTAKEAKEYFA\D
		1	l	I		MERHRILFRYAGPEDDAAITLA
		1		l		
1		1		I		FSKKKIDDRKEWLTNFMEDRR
				I		QRRLHGLPEQFLYGTATKHLTY
1		1	1	I		NDFINKELILFSNSDNERSIPSLV
22222	******	ļ.	00410	0.7	200	DGFKPGQRKV
22389	52757	A	22517	87	292 903	
22390	52758	A	22518	1		
22391	52759	A	22519	3	311	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, -possible nucleotide insertion)
				sequence	1	
00000	50750	<u>_</u>	100500		1,000	LANGE AND ADDRESS OF THE PARK OF
22392	52760	Α	22520	l .	1087	MPSAAWKRPVEGPATLRRPLGI
					1	AAASQCPSSPQPGCLIGAVNLK
					i	SSNRTPVVQEFESVELSCIITDS
						QTSDPRIEWKKIQDEQTTYVFF
					İ	DNKIQGDLAGRAEILGKTSLKI
			ľ		i	WNVTRRDSALYRCEVVARNDR
						KEIDEIAIEVT\VQVKPVTPVCR
						VPKAVPVGKMATLHCQGE*GP
			ľ		i	PPA\HYSWYRNDVPLPTDSRAN
				1	ŀ	PRFRNSSFHLNSETGTLVFTAV
					İ	HKDDSGQYYCIASNDAGSARC
						EEQEMEVCHTDKRGGFPWSW
						AAPHLWFLQDTALLPAALTGW
						CSVSVAFPGAQCKLSVDLSFWG
		ŀ				LEDGGPLLTAPLGSTPVGTLCR
						GSDPTFPFCTALADDLYESPTPA
						ANFCLDIQAFSYIL
22393	52761	Α	22521	628	1117	FFISVINGQVSSVQRLSGVGPAC
						LSCGSANPGPPPGTSPGAGAQR
						R*\PRADGSGSPQWPRGARVGG
				ļ		GRLGTGGRGRPGWROVPRRLS
						PGFGR*GGTGPGPVGTSGKRGP
						SRRRAPANDKAACWPRFPGOP
						AS*TGFRGERGVKGFSSWGSG
						WRAWEDGGTVH
22394	52762	A	22522	2	238	
22395	52763	A	22523	131	2891	
22396	52764	A	22524	131	3103	LVSFLLFQDALPEGDASPLGPY
						LLPSGAPERGSPGKEHPEERVV
						TAPPSSSQSAEVLGELVLDGTA
						PSAHHDIPALSPLLPEEARPKHA
						LPPKKKLPSLKOVNSARKOLRP
						KATSAATVQRAGSQPASQGLD
						LLSSSTEKPGPPGDPDPIVASEE
						ASEVPLWLDRKESAVPTTPAPL
l			1		1	OISPFTSOPYVAHTLPORPEPGE
l					1	PGPDMAQEAPQEDTSPMALMD
1		l				KGENELTGSASEESQETTTSTIIT
1			1		1	TTVITTEQAP
	l	L_	l	l		ITTYTITEQAF .

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22397	52765	A	222525	131	2527	SASPLGPYLLPSGAPERGSPGKE HPEREXVTAPPSSSOSAEVLGE LVLDGTAPSAHDIPALSPLLPE EARPKHALPPKKKLPSLKQVNS ARKQLRPKATISAATVQRAGSQ PASQGLDLLSSSTEKPGPFGOPD PIVASEEASEVPLWLDRKESAV PITTAPLQISPFTSQPYVAHTLP QREGEGFPDMAQEAPGEDTS PMALMDKGENELTGSASEESQ ETTTSTHITTIVITEQAPGMC THOMOGORY THOM
						LLPSGAPERGSPÖKEHPEER VV TAPPSSSQSAEVLGELVLDGTA PSAHHDIPALSPLLPEEARPKHA LPPKKKLPSLKQVNSARKQLRP KATSAAT VQRAGSQPASQGLD LLSSSTEKPGPPGDPDPIVASE ASEVPLWLDKKESAVPTTPAPL USPFTSGPVYAHTLPORPEPGE PGPDMAQEAPQEDTSPMALMD KGENELTGSASESQETTTSTIIT TTVITTEQAP
22399	52767	c	22527	161	352	
22400	52768	A	22528	2	368	LGSTHASVRINILSVDVLYFIFY RRDDNKDIDSEKEAAMEAEIKA ARERAIVPLEARMKQFKDMLL ERGVRKQSLGEIFLFCISNINMS C*KVKSKVSAFSTWEKELHKIV FDPRYLLLNLI

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22401	52769	IA	22529	2	614	KDHRWESGSLLEREEKEKLFNE
22401	32709	ľ	22329	2	017	HIEALTKKKREHFROLLDETSAL
		1				TLTSTWKEVKKIIKEDPRCIKFS
		1				SSDRKKOREFEEYIRDKYITAK
		1	1			ADFRTLLKETKFITYS*FSERCE
			1		i	SESL*KMLGLAAPQTEVLCSST
ļ						
		1			i	E\RSKKLIQESDQHLKDVEKILQ
						NDKRYLVLDCVPEERRKLIVAY
		_				VDDLDRRGPPPPPTASEPTRRST
22402	52770	В	22530	201	296 1290	
22403	52771	В	22531	1	3369	OCCUPALED CONCEPTED THE
22404	52772	A	22532	3	3369	SSSVMAERGGVGGESERFNPGE
						\LRMAQQQALRFRGPAPPP\NA
ĺ					1	AMRGPPPLMRPPPPFGMMRGPF
						PPPRPPFGRPPFDPNMPPMPPPG
						GIPPPMGPPHLQRPPFMPPPMSS
						MPPPPGMMFPPGMPPVTAPGTP
						ALPPTEEIWVENKTPDGKVYYY
					l .	NARTRESAWTKPDGVKVIQQS
1						ELTPMLAAQAQVQAQAQAQA
						QAQAQAQAQAQAQAQAQA
		1	l			QAQAQAQAQAQAQAQAQA
		1			1	QAQAQAQAQVQAQVQA
22405	52773	A	22533	43	347	
22406	52774	Ä	22534	1	1776	
22407	52775	Ā	22535	3	433	PLAPNHLGGGAVVLGAESHAS
		Γ.		l	1	KDVAIDMMDSRTSQQLQLIDE
		1				QDSYIQSRADTMQNIESTIVELG
		1				SIFQQLAHMVKEQEETIQR*DR
		1				VNSAHPWDPVHLLPSTOOVFIE
		ı				HLFM
22408	52776	A	22536	304	1403	SRGNATGLRTRIRVSTWVSQRH
22400	22770	ľ,	22550	150.	1	RSCPLQLLAVAAATSSPLPPPVT
l		ı				LVPPPPDTMSCRDRTOEFLSAC
		1			i	KFAARPVKNGI\QTN*GQLLRW
		ı				LVRQRSEIHPSLAKRHLGKDLS
		ı				NTFAKLEKLTILAKR\KPLFDDK
		ı				AGEIEELTYIKODINSLIKOIAO
		ı				LO\DFVRAKG\SQSGRHLQTHS
		ı			1	NTIV\VSLQSKL\ASMSNDFKSV
	1	1				
		1			1	L\EVRDRGTLKQ\QREPEERQFS
		1			1	RAPV\SALPLSPTHRAGGA\VVL
		1		1	1	GAESHASKDV\AIDMM\DSRTS
		ı		1		QQLQLIDEQDSYIQSRADTMQN
	l	1				IESTIV\ELGSIFQQLAHMVKEQ/
		ı			l	ERETIQRIDENVLGAQLDVE\AA
		1				HSEILKYFQSVTSNRWLMDKIF
22409	52777	В	22537	147	506	LILIVFFIIFVVFLA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22410	52778	A	22538	2	1397	AAPGNGLSPSSWRSELACEFOH
		l				ECTAEVRFGAVVCRGARSHSFY
					i	LEREILQ/PPGMSL/PENPVTPML
	1				l	RHLMYKIKSTGPITVAEYMKEV
	1		ł			LTNPAKGYYVYRDMLGEKGDF
	1					ITSPEISQIFGELLGIWFISEWMA
		ł				TGKSTAFQLVELGPGRGTLVGD
		1				ILRVFTQLGSVLKNCDISVHLVE
		l				VSQKLSEIQALTLTKEKVPLER
	1	l		l		NAGSPVYMKGVTKSGIPISW*S
		l				RICTD\VPKGYSFYLAHEF\LDV
		1	1			LPVHKFQKTPQGWREVFVDIDP
						QVSDKLRFVLAPSATPAEAFIQ
1						HDETRDHVEVCPDAGVIIEELS
	1		1			QRIALTGGAALVADYGHDGTK
	İ					TDTFRGFCDHKLHDVLIAPG\T
	į.	l	l			ADLTADVDFSYLRRMAQGKVA
	į.					SLGPIKQHTFLKNMGIDVR/L*R
	1					FF*INQMRPSVRQQLLQGYDML
						MNPKKMGERFNFFALLPHQRL
	t				i	QGGRYQRNARQSKPFASVVAG
						FSELAWQ
22411	52779	Α	22539	2	377	
22412	52780	Α	22540	2	1427	
22413	52781	Α	22541	2	568	CSSPSN/TILGVQKLNSQWRLA
		l				QDFRLINEAVIPLYPVVPNPYTL
		[				LSQIPEEAESFTVLDLKDAFFCIP
		l			i	LHSDSQFLFAFEDPTDHNSQLT
		l				WMVLPQGFRDSPHLFGQAQAQ
		1				DLVHFSSPGTLVVQYMDDLLL
		l				ATSSEASCQQATLDLLNFLANQ
		1		1		GCKRVGIALGVLTQTHGTTPQP
l	1	ł	1	l	i	VAYLSKETDVAAK

SEQ ID   SEQ ID NO:   Met   SEQ ID NO:   Not	Amino acid sequence ( X=Unknown,
	*-Stop codon, /-possible nucleotide
pequence   109/540,217   readon for peptide joi peptide sequence	deletion, \=possible nucleotide insertion)
sequence	
22414   52782   A   22542   I   1230	GPRGTGPKGKARSEKGCSLSHG
22414   52782   A   22542   I   1230	
	PQTNKPLVVQKGQKMEQANHP
	VGLVISVVYKDILKKIVQRETS
	HPLIHVRYAEAITGRRTAPEDK
	GSLGRDMLAKAGAIIYMNMGN
	KLPIWCHLLEEGIYLEVWALEG
	QFGRAKNACPVQIRLKDPTTFP
	YQRQYPLRPEAHKGLQDIVKH
	VKAQGLVKKCSSPCNTPILGVQ
	KPNGQWSLVQDLRLISEAVIPL
	YPVVPNPYTLLSQIPEEAEWFP
	VLDLKDAFFCIPLHYDSHDSQF
	LFAFEDPTDHTSQLIWTVLPQG
	FRDSPHLFGQALAQDLGHFSSP
	GTLVLQYVDDLLLATSSEASCQ
	QATLDLLNFLANQGYKASRSK
	AQLCLQQVKYLGLILARGTRTL
	GKERIQPILAYPHPKTLKQLWG
	FLQIT/GFCQLWIPR*SKI
22415 52783 A 22543 1 211	
22416 52784 A 22544 1 472	
22417 52785 A 22545 1 393	
22418 52786 A 22546 59 411	SWPSDKQTLVVQRGQKMEQA
	NHPDPTDHMSQLMWT/VLPQG
	FRDSPHLFGQALAQDLGHFSSP
	GTLVLQYVDDLLLATSSEASCQ
	QATLALLNFLANQGYK/LSRSK
	AQLCLQQVKYLCL
22419 52787 A 22547 43 236	TSSLLRKEDSAPSQGKSVVFTL
	TSQG*VPLWRTLQLQGPFFTPSS
	RK*LELSSANSQQHLGCPV
22420 52788 A 22548 1 1446	
22421 52789 A 22549 I 1521 22422 52790 A 22550 2 689	
	GUDDTEDD COLUMN TON
22424 52792 A 22552 636 1586	SNDRTEDDCGKHPFMSSPP\TEP
	WVCLIEGQEIDFLLDTGTTFSV
	LIPCLGRLSSRSVTIQGILGQPVT
	RYFSHLLSCNWETLLFSHAFLV
	MPESPTPLLGRDILAKAGAIISM
	KTGNKLPICCPLLEGINPEVWA
	LEGQFGRAKNAHPLQIRLKDPI
	SFPYQRQYPLRPEAHKGLQDIV
	KHLKAQDSVRKCSSPCNTPILG
	VQKLNSQWRLVQDLRLINEAVI
	PLYPVVRNPYTLLSQVPEEAEW
	FTVLDLKDAFFCIPLHSDSQFLF
	AFEDPTDHTSQLMWMVLPQRF
	RDSPHLFGQAQAQDLGHFSSPG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		Ļ.	00550	292	1.100	
22425	52793	A	22553	292	1493	SNDRTEDDCGKHPFMSSPP\TEP
						WVCLIEGQEIDFLLDTGTTFSV
		1				LIPCLGRLSSRSVTIQGILGQPVT
1		1				RYFSHLLSCNWETLLFSHAFLV
						MPESPTPLLGRDILAKAGAIISM
			1			KTGNKLPICCPLLEGINPEVWA
		1				LEGQFGRAKNAHPLQIRLKDPI
		l	1			SFPYQRQYPLRPEAHKGLQDIV
		İ	i			KHLKAQDSVRKCSSPCNTPILG
		ŀ			\	VQKLNSQWRLVQDLRLINEAVI
		1				PLYPVVRNPYTLLSQVPEEAEW
		l				FTVLDLKDAFFCIPLHSDSQFLF
						AFEDPTDHTSQLMWMVLPQRF
1		١				RDSPHLFGQAQAQDLGHFSSPG
		l		l		TLVLQYSEIAKTLYTLIKEMER
]		1				ANTHLVEWEPEAETAFETLKQ
		1				ALVQAPALSLPTGQNFALYVIE
		1				RAGIALGVLTQTHRTTPQPVAY
22426	52794	Α	22554	2	307	
22427	52795	Α	22555	3	400	
22428	52796	Α	22556	2	78	
22429	52797	Α	22557	395	1336	SNNRTEGASSCH\PLTEPQVCLT
İ		l				IEGQEIDFLLDTGTAFSVLISCPA
		l		l		WGTGQDSVSKKNEKKRKEKA
		l				KGNHQANAEAKIAARWNPSLE
		ı		i		IPTEGPLVWNNPLQEIKPQYSLT
1		l				ETEWGLSQGHSFLPLGWLMTE
		l		İ		EVKVLIPEASQWKILKTLHQTF
		l		1		QMGIENTQQMAKSLFTGPNLL
		l				WTIRHVVKACEVCQRNNPLVH
		1				HKAPLGEQRIGHYPGEDWQLD
1		1				FTHMPKSKGFQYLLVCVDTFT
		ı	l			NWIEAFPCKIEKAQEVNKVLIH
		1	l			ETIPRFGLPQSLQSDNDPAFKA
		l				MITQGIFQGAKDTISPSLCLEATI
		1				LREGREGN
22430	52798	Α	22558	244	638	GPLNPTFLGPPCGLGGKLVFLL
		l				LRQ*VQLFRSAGSRDRCGFLDR
		1				ERNKTNQNCGKQAKKSPKDHK
		1				NPRAVGYVPFNL*QEGNLAQP
				1		GYMSPSPSLI*SRSRQTWGSFQ
		_	l .			MILIGTKRNRPKRKSEIRERLQP
22431	52799	Α	22559	210	600	RGKRTWRERRNGWSRGFCPLP
	1	1	l			LTGSWLISRESPRLACLGEGLG
						VPGM*DLPHAWPTTLMCPELV
			1			GSWSH*LQE*SHGPSQ*VLQLL
			ŀ	l	i	RWCVWSLFLLMFGCVWSFFLL
		1	1			VSSWSRWLRSEATDLCGECYSS

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nueleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22432	52800	A	22560	2746	5211	ETSCPLVRVHYAEAITGRCTAP
						EDEGSLGOKPPARRSNNRTEGA
					i	WGKRQLMSSP/STEPRVCLTIEG
l					İ	QEIDFLLDTGAAFSVLISCPGRL
						SSTSVTIQGILGQPVTRYFSHLL
	ŀ					SCNWETLLFSHAFLLMPERPTG
			l			LLGRDILAKAGAIIYMNMGNKL
1						PICCP\LLVEGINPEVWALEGOF
l	Ì					GRAKNARPLQIRLKDPTTFPYO
						ROYPLRPEAHKGLODIVKHVK
1						AQGLVKKCSSPCNTPILGVQKP
Į	į	ĺ				NGOWSLVODLRLISEAVIPLYP
						VVPNPYTLLSQIPEEAEWSWAL
						TFKDAFFCIPLHSDSQFLFAFED
				İ		PTDHTSQLTGTVLPQGFRDSPH
						LFGQALAQDLGHFSNPGTLVLC
						YVDDLLLATSSEASCEQATPDL
						LNFLTNQGYKVSRLKAQLCLQ
	Ì		l		ļ	EVKYLGLILARGTRALSKE*IQP
	ľ				ł .	ILAYPHPKTLKOLRGFLEITGFC
١.	ł					*LWIPGYSKIARPLYTLIKETQR
	1					ANTHLVEWEPEAETTFKTLKO
l						ALVOAPALSLPTGONFSLYVRE
	l					RARIALGVLTQTHGTTPQPVAY
						LSKEIDVVAKGWPHCLRVVAA
						VAVLASEAIKIIOGKDLTV*TTY
1					l	DVNGILGAKGSLWLSDNCLLR
						YQALLLEGPVLQICTCMALNPA
						TFLPEDGEPIKHDCQQIIVQTYA
						ARDDLLEVPLTNPDLNLYTDGS
						SFVENGI/RKVSDVTILESKPLPP
						GTSAQLAELVALTWALELGKG
22433	52801	A	22561	903	4440	
22455	32801	A	22361	903	4440	DHPGHIHSSPCHERPNNVSTLPF
		ĺ				LPGEQHPILLPPRNCPGDKILEE
						NFRYNNYKRTMMSFKERLENT
			ŀ			VERCAHINGNRPRQSRGFGELL
						STAKQDLVLEEQSPSSSNSLENS
						LVKDYIHYNGDFNAKSVNGCV
		1	1			PSPSDAKSISSEDDLRNPDSPSS
			1			NELIHYRPRTFNVGDLVWGQIK
	1		1			GLTSWPGKLVREDDVHNSCQQ
						SPEEGKVEPEKLKTLTEGLEAY
						SRVRKRNRNGIKVPPRSPRSTIG
		L_				SPRPSMPSSPS
22434	52802	Α	22562	I	501	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22435	52803	A	22563	1688	2038	I SATHGLSDSCYWLVVSRAIGRN
22433	32003	^	22303	1000	2038	NLSPASPASIIHALVFLAEPOLF
						OPRILGLIDEESGHEGGRPAAKT
	1		ļ			RKLSGPWSRPAAFMSTLLINOP
	1				i	QYAWLKELGLREENEGVYNGS
						WGGRGEVITTYCPANNEPIARV
						ROA\SVADYEETVKKAREAWKI
	1	ĺ	l			
	1					WADIP\APKPKEKLVRQIG\DAL
						REEDSKLLRKPWCPLEDWGKIL
			i			LEGVGEVQEYVDICDYAVGLS
						RMIGGPILPSERSGHALIEQWNP
						VGLVGIITAFNFPVAVYGWNN
		1				AIAMICGNVCLWKGAPTTSLIS
						VAVTKIIAKVLE\DNK\LPGAICS
						LTCGGADIGTAMAKDERVNLL
	i					SFTGSTQVGKQVGLMVQERFG\
		l				RHYQNWRSLLELGGNNAIIAFE
	1	1				DADLSLVVPSALFACCGEQLGQ
						RCTT/SRGRLFYYMESIHDEVFK
					1	QTLKRAYGTRSRVG\NPW\DPN
	1					VLYGAISHQGRQLSMFLGAVEE
	1					AKKERLAQWVYGGKVMDRPG
	1					N\YVEPTNVTS\LDHDASIAHTE
	1					TFAP\ILYVFK\FKNEEEVFAWE*
	1					WK*NRGLSS*HSLPKDLGAESF
	1				1	RW\LGP\KGTQTCG\IVKCQPFPT
			ĺ			KWGLRLGGA\FGGEKAHLVVA
			l			RESGQ*C/WGKQYMRKVYLVL
			l			FNYS\KDLPSGPQG\IKFQ
22436	52804	В	22564	1	1434	
22437	52805	В	22565	1	1185	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22438	52806	A	22566	1	2450	GQACHASSSPLKGSRGSPPNFEI
		1				LYGPIFEDSLAPD*PKVIAGQ\IT
	1	1				LRDAVTGPISSPNEAGKIYOLKE
		1		1		NPAVLICRVRGLHLPEKHVTW
					ŀ	RGEAIPGSLFDFALYFFHNYOA
1					1	LLAKGSGPYFYLPKTQSWQEA
					İ	AWWSEVFSYAEDRFNLPRGTIK
		ı				ATLLIETLPAVFOMDEILHALR
					1	DHIVGLNCGRWDYIFSYIKTLK
						NYPDRVLPDRQAVTMDKPFLN
						AYSRLLIKTCHKRGAFAMGGM
i		ı				AAFIPSKDEEHNNQVLNKVKA
						DKSLEANNGHDGTWIAHPGLA
		ŀ				DTAMAVFNDILGSRKNQLEVM
						REQHAPITADQLLAPCDGERTE
1						RKFNALMESEKGQSQNLWKFA
						VYSGLRHGELAALAWEDVDLE
				1		KGIVNVRRNLTILDMFGPPKTN
						AGIRTYTLLQPALEALKEQYKL
1						TGHHRKSEITFYHREYGRTEKQ
i						KLHFVFMPRVCNGKQKPYYSV
		1				SSLGARWNAAVKRAGIRRRNP
		1				YHTRHTFACWLLTAGANPAFIA
						SQMGHETAQMVYEIYGMWID
						DMSDEQIAMLNARVMVYDVA
1					l	SGNALFISELGPLPENVTWLSPE
				1	l	GEFOKWNGTAWVKDTEAEKL
				1	l .	FRIREAEETKNNLMQVASEHIA
	ł	i				PLQDAADLEIATEEEISLLEAW
						KKYRSIPLCSKVVTMMFDDRP
l	ŀ					GCPVGFPMWGRGGFDRMPPG
		1		1		WGGRPMPPSRRDMSPHRGPPP
22439	52807	A	22567	1	941	MIPNVOGPMRGLEGPNYIFIKIS
22439	32807	l <sup>A</sup>	22307	l'	941	KNRRYTMAVTSVGLASSLCRV
1					1	VRPCRAPAGVVSRGSPDDEWL
1					l	
ĺ						FCPQARIQQQQDID/NGTLPDFIS ETASIRDA/DWKIRGIPADLEDR
						RV/EITGPVERKMVINALNAI\F
	I					
	1		l		į .	MADFEDSLAPDW/NKVIDGQIN LRDAVNG/TISYTNEAGKIYQL
					l	
1		1				KPN/PA VLICRVRGLHLPEKH/V
1		1				TWRGEAIPGSLFDLA/LYFFHNY
1	1	1	1	1	I	QALLAKGSGP/YFYLPKTQSWQ
1		1				EAAWWS/EVFSYAEDRFNLPRO
1	1	1	1		1	T/IKATLLD*TLAGGFQMVDPSA
				1		LGG*TGPWDHLRYIKVENIPKV GEEGDAETTGGTAV
22440	52808	A	22568	2	393	GERODAETTOGTAV
22441	52809	Â	22569	1066	1824	
22442	52810	A	22570	2	165	
22443	52811	A	22571	3	902	
22773	102011	٠,,	1		P	

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22444	52812	A	22572	1	2034	
22445	52813	A	22573	176	873	RLFGEMSKLN/HNLSEVMSKLE
	1					KQHSSKVFVVKRLSSSSRRSLVI
	1					SPPVRTSTVSSPLTSPTSPSTLSS
	i				1	KSKSESVSATEDLAPDAAQGED
	1					NSEIKELLEDEEIGKEGSEASSS
	1					EEEEPLLTCNGPSQAQPSPIIESA
		ŀ				RSQEEVPPSSPALSPGGALSPSG
	1				l	HPLSSAAEVVLRTRTAVPQAKG
i	1					NNVVITSYMTNRGFFEDKKATF
	İ					APSFLMNIKGNKTSVVKNSILE
						QGQLTVN
22446	52814	A	22574	1	1890	
22447	52815	A	22575	2	1279	LSVFPRVRNFTTQAECH/KLFK
	1				ļ.	VLKNFLIAGGGVHESSKRAS*T
						LQFIYY\SEWYGHEELKAIVWN
						NDLLWEDYEEKLADQAVRTM
						EIYVAQFSEIKERIAKRGRKLVD
						YDSARHHLEAVQNATKKDEAK
		ĺ				TAKAEEEFNKAQTVFEDLNQEL
	1					LEELPILYNSRIGCYVTIFQNISN
	ŀ				1	LRDVFYREMSKLNHNLYEVMS
		l				KLEKQHSNKVFVVKGLSSPSTL
		1				SLKSESESVSATEDLAPDAAQG
		l				EDNSEIKELLEEEEIEKEGSEASS
		1				SEEDEPLPACNGPAQAQPSPTT
	l	l				ERAKSQEEVLPSSTTPSPGGALS
1					1	PSGQPSSSATEVVLRTRTASEGS
		1				EQPKKRASIQRTSAPPKPPEKPV
1	l	1			l	RTPEAKENENIHNQNPEELCTSP
		1				TLMTSQVASEPGEAKKMEDKE
	l	1			<u> </u>	KDNKLISANSSEKYIFQRPSVLI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22448	52816	A	22576	1	1791	MPGARTSSSGASENHRARGOG
						GGPQGVGRMAEGKAGGAAGL
		l				FAKQVOKKFSRAQEKAL*KLG
	١.	l			1	KAVETKDERFEQSASNFYQQQ
		ļ	l			AEGHKLYKDLKNFLSAVKVMF
		1				ESSKRVSETLQEIYSSEWDGHE
		i				ELKAIVWNNDLLWEDYEEKL\
						ADQAVRTMEIYVAQFSEIKE\RI
						AKRGRKLVDYDSARHHLEAVO
		l				NAKKKDEAKTAKAEEEFNKAO
		1				TVFEDLNQELLEELPILYNSRIG
						CYVTIFONISNLRDVFYREMSK
			İ			LNHNLYEVMSKLEKOHSNKVF
			İ			VVKGLSSSSRRSLVISPPVRTAT
		1		1		VSSPLTSPTSPSTLSLKSESESVS
						ATEDLAPDAAQGEDNSEIKELL
						EEEEIEKEGSEASSSEEDEPLPA
						CNGPAQAQPSPTTERAK\SQEE
		l				VLPSSTTPSPGGALSPSGQLSSS
						ATEVVLRTRTASEGSE\RPKKT
						ASIQRTSAPPRRPPPPRATASPRP
						CSGNIPSSPTASGGGSA\TSPRAS
						LGTGTASPRTSLEI\*PNPE\PPEK
						PVRTPEAKENENIHNQNPEELC
			ľ			TSPTLMTSQVASEPGEAKKME
			ł	İ		DKEKDNKLISADSSEGQDQLQV
						SMVPENNNLTAPEPQEEVSTSE
22449	52817	Α	22577	2216	2568	RPSLGLGM/C/TIWQ/TSLA/GKH
						RRTIETHTDDYRISLEQGTPGLE
			1			PPLNALTAISKKIGCNGFFPFQIR
						PGKNETDGRMFSPAIGIVEDPV
						TGNANGPMGAWLVHHNVLPH
						DGNVLRVKGH
22450	52818	Α	22578	108	532	WGSKGNLLYSHQTHSVSHQFS
						LVACCLELEYSPFSFPYACLLSS
						LMRLEKHQELNLWPAHQRSMR
						HPQDTLVVFS/HHGRVLL*DKT
	l					SRGTIRQQHSRFTKICCSAATAA
						VTQANRVWSGPLANSNRPAAE
						GPVLAAFMGWL
22451	52819	A	22579	1	1125	

SEQ ID	SEQ ID NO:					Amino acid sequence ( X=Unknown,
NO:	of peptide	hed	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22452	52820	A	22580	1	985	MCARGSLWGCYSGLERKCEAA
		l				LPDRVVSSKSACGAVSQDLLQ
		1				AOGRRAGPPSWLYADPVHTGT
						IECGGHWCLNAALLDFQSPQRL
		ŀ				LTYLHCGLQRPMESDPPELAPM
						VHHQEGIYLNIRKAMYDKPTA
					i .	NKLNWDKSNVSSLITGTRQEET
		1				TANGTTFATIHLQFFIQIPAVPR
ŀ						KTLLNRADYYFPEHNPSLTLVI
						NRRITKFSEPNWEAPPSRGRLTP
		1				HTAGYSSETKIPEERSDSSIRGS
		l	l			RKSAVLQPLLLDSQANRVWSG
		1				PLANSNRPAAEGPVC*LWWVT
						LHGPLKATMQVCQQPLW
22453	52821	Α	22581	294	934	YPPSPAELTVCRLRQLQLRDVL
		l				YCIAYPSGPGVKPFRGNSA/G/V
						VFPADNLSEAQMQLI/ARELGH
1		l				SETAFLLHSD/DSDVRI/RFTPTV
		l				EVPI/CGHATVAAHYVRAKVL/
		l				GLGNCTIWQTSLAGKHR/VTIE
		l				KHNDDYRISLNALTAIS/KKIGC
		l				NGFFPFQIRPG/KNETDGRMFSP
		l				AIGIVE/DPVTGNANGPMGAWL
22454	52822	В	22582	1163	5585	VH/HNVLPHDGNVLRVRQ*GPR
22454	52822	A	22582	2	429	AACAAAMSLVIPEKFOHILRVL
22433	32023	<u>ا</u> ا	22363	1	1427	NTNIDGRRKIAFAITASSSSSSS
ŀ		1				SSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSS
l		1				SSSSSSSSVNLLSNGLYLVKG
		1				VGRRYAHVVLRKADIDLTKRA
		1			i	GELTEDEVERVITIMONPROYKI
		1	,		l .	PDWFL
22456	52824	A	22584	84	424	GVHAASDKPCWITTSQTPDVQ
		l				HRW/WLLISSML*TAPNRRSDK
		1			l .	AFTPHPTNHAGSQLRKHPTCSI
		1			l	DGVIDIFHAVNTFVQHHMEETL
		l				KAFHVTGHHILIARHGLGVGKE
		l				NTEHAANVVNDQRNSGFFCRF
		1				QQTFCQTCCQLIQRFVNARLRD
		l				FISC
22457	52825	Α	22585	88	381	IQVLANGLDNKLREDLERLKKI
		l			\	RAHRGLRHFWG*VGGLISLPTS
		l				TQHSSYSLFFFPNLLFLLCMTCD
1						SSLFTCSLRVRGQHTKTTGRRG
		L				RTVGVSKKK
22458	52826	A	22586	355	699	FIDTTEAGSHFLYALVRFAATIT
						TITDDGARRLRTLFQRQQADLV
		1	[			SVSKSGFFTADCTHTNALIDVV
		1		1		RTIFNNAVF*NPGFVIARLKIEIA
			į.			IIKATFGQLTEDREQVLMFQAV
l		_				RR

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide scquence	deletion, \=possible nucleotide insertion)
22459	52827	Α	22587	1389	1532	SVLPGRPDARQYRQTRQSW*R WSLRLRGSYPVAGRSLSRRHRQ TERF
22460	52828	Α	22588	1	427	LVHQQWIQTKKKSLIYPFCHSP SQRVVEVAALDQWLAMLYPR VEHQAVLAVVVVVVGPLVVRP YHTRRDAVRLLRQPAVFIVAV DIVRILRDAVIRPHCECTVRVIR SGSGSRLPAGSVAYGMRMSRS GARVRTLSILPVAL
22461	52829	Α	22589	I	915	
22462	52830	Α	22590	1166	1321	
22463	52831	Α	22591	813	977	
22464	52832	A	22592	3485	4237	
22465	52833	A	22593		499	ARAAACAAAMSLVIPEKFÖHLL RIVLNTNIÖGRRKIAFAITAIKG VGRRYAHVVLRKADIDLTKRA GELTEDEVERVITIMONPTQYK RIPDWFLNROKDVKDGKFQFGS *ANGLSTTKLIKENLKPLKEDSG PIKGLRQLPWGLRVRGQHTKT TGRRGRTVGVSKKK
22466	52834	Α	22594	1	1197	
22467	52835	Α	22595	I	1905	
22468	52836	A	22596	37	514	WTWWLRCYENRHRHCHLLQQ LPTSPHQHAVATPTTVAVPTAP AATPPVVVAVVVAPVVVVVVP VVPVVASVPFVEPGSDVIVCAE MDEQWGYVGAKSRQRWLFYA YDSLRKTVVAHVFGERTFDVVI WMTDGWPLYESRPAFTGRDC VDDAFFAVAGV
22469	52837	Α	22597	1	471	
22470	52838	Α	22598	2	323	SRIQPGSDVIVCAEMDEQWGY VGAKSRQRWLFYAYDSLRKTV VAHV/FGERTIRTLTRRLFRCLR SSHDLVSSPVSHRRRRAGSRQR HRRARGGENVEYQWGALRHG
22471	52839	Α	22599	2033	2310	ARYVGNAGCTGGALFGRSQTI REAVG/ERRLAESRDVAALDTP CDPEIKMAEWMQTLKETGSTS GHIVTRRISVRISHVTPGPASQD GPDVQQR
22472	52840	В	22600	I	4891	
22473	52841	В	22601	7	1233	
22474	52842	Α	22602	193	492	
22475	52843	Α	22603	1	486	L

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22476	52844	A	22604	62	443	RSDRLLIHGFSFPAMGFRLRKT VVAHVFGERTMATLGRLMSLL SPFDVVIWMTDGWPLYESRLK GKLHVISKRYNTORIERHNLNLR OHLARLGRKSLSFSKSGTACCG
22477	52845	A	22605	971	1444	HCLTROIPMLHHQGNQLF SMGSGDMRSSPSRAAALRQLPR PFRQLAVRMEMISRFRSAKEQT KSLRKWRKAKSIF*SVAQTGFN VLVCTTIIETGIDSRQPTLLSLNA RITSVCAAAPVTRSRTFASSGY AWLLTPHPKAMTTDAKNVLKQ
22478	52846	A	22606	1	1047	LPRWKISGQVRAGTARSGDSRR W
22479	52847	A	22607	318	598	LTGFRRLCDLRFGGGAVRSVLA RSREFILQH*FRRIVTGSMGRSL VPSAPLR/PSIVLVLFRKAW*LV KTQRKMHVVSIVQKKASCWYP LLGCL
22480	52848	Α	22608	1	2652	
22481	52849	Α	22609	1	852	
22482	52850	Α	22610	1	807	
22483	52851	A	22611	1	1740	
22484	52852	A	22612	I	3144	

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ĺ		sequence		
22485	52853	Α	22613	173	1983	PVQNGSSPLGALPGAPLIFGPSV
						PAISRPGSCSPEWPTLDTSGYSA
						PR/WLFTAYAGSEAVFAHVFGD
						ALGDAGLLMSLLSPFDVVIWM
				l		TDGWPLYESRLKGKLHVISKRY
						TORIERHNLNLROHLARLGRKS
1				Į.		LSFSKSVEOHDKLESLPGOIDKS
						LDEASLSLRAGSLRTITHILLPLL
						RPAILSALIYSFVRAITTVSAIVF
						LVTPDTRVATAYILNRVEDGEY
		l		i		GVAIAYGSILIVVMLAIIFIFDWL
						IGESLLSRFEKTVDDHSQSRLPA
1						EGSILSQDISGVKLFSLLTPARG
1						YVPIRAPCPFLCASYRFFCVARQ
1			ŀ	ŀ		QGLDAALIDAPLAVDALTPLSE
				i	l	LLSHVGQAPCSVPVVDEDQQY
				l		VGIISKGMLLRALDREGGVRVP
			ĺ		1	VDYILNGFHPLLLGMPAPLAIIV
		l				FALIACRFPGSNGRIRRSRRIRH
		l				LHKMPDATLSASYQAYVRHQT
1						NVTALVLRSSHRFRKDRHCWG
1	ļ.	l				FVFLQQPILRATGQFEIQTFRYL
1				1		SVFSVQRHKTNGKTALMSGNV
						ADKAVNRATGPTACVWLFSLR
	l	ı	l			NSSMVTTRFCGLCEWADOOGA
	İ				1	RPARISENGKRGYSYLYDVHM
		)				GAQSGETRGRQGEYASKPPSRE
22.406	52054	l .	22614		2149	GAQSGETRGRQGETASKFFSRE
22486	52854	В	22614	I.		
22487	52855	A	22615	1	1815	
22488	52856	Α	22616	2014	3411	E A LA COLLO DEL LA COLLO DE L
22489	52857	Α	22617	1	922	MVICQPINGGWQWPPSTGTDP
	l	1	i			APYFRIFNPTTQGEKFDHEGEFI
	l		1		1	RQWLPELRDVPGKVVHEPWK
1		ı			1	WAQKAAAPPQTRLHSTPTTIPH
1		l			i	RVPGHTPNLKPIWEVDLPKFSH
		1	1		1	KATSHATDKPSDTTQNDQIVIA
1		i i				YMEPPEIVHETTTSSTHPSLYIIL
		l				KRQTRELRPQSVTSRIQPGSDVI
1		l				VCAEMDEQWGYVGAKSRQRW
		l	1			LFYAYDSLRKTVVAHVFGERT
		l		ı		MATLGRLMSLLSPFDVVIWMT
		İ				DGWPLYESRLKGKLHVISKRYT
1		1				ORIERHNLNLROHLARLGRKSL
1	l	ı		1		SFSKSVELHDKVIGH\YLNIKHY
L	l					DI DIO VELLIDIA VIOTA LINIATI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22490	52858	Α	22618	1458	2185	YGHEWRWMPGNRPH\YGRWP
		l			i .	QHDFPPFKKLRPQSVTSRIQPGS
		l			i	DVIVCAEMDEQWGYVGAKSR
	İ	l		l		QHWLFYAYDRLRKTVVAHVF\
	ŀ	1				GERTMATLGRLMSLLSPFDVVI
	ľ	l		l		WMTDGWPLYESRLKGKLHVIS
		l		l		KRYTORIERHNLNLROHLARLG
		l				RKSLSFSKSVEKITSGPTTNATP
		l		1		GTOKTATDKRGGKKLPIQIPIGP
		l		ļ		ISRIASGAAINTISIGLRKFLVTA
		ı				GVMLSTQRSIYDRHQVITSAGIT
22491	52859	A	22619	493	1518	
22492	52860	Α	22620	963	1597	
22493	52861	Α	22621	1	3052	MATDSVGLNCTPAPHARDVLL
1		l				WRPSSLPVSTSCVFPLGRIGDVS
	1	ı		ì		DESDDKTEAPTPHRLEKAREEG
		ı				QIPRSRELTSLLILLVGVSVIWF
		1	İ			GGVSLARRLSGMLSAGLHFDH
		1				SIINDPNLILGQIILLIREAMLAL
		1			l	LPLISGVVLVALISPVMLGGLVF
		1				SGKSLQPKFSKLNPLPGIKRMFS
		l				AQTGAELLKAILKTILVGSVTG
		l			ĺ	FFLWHHWPQMMRLMAESPITA
					l	MGNAMDLVGLCALLVVLGVIP
						MVGFDVFFQIF
22494	52862	Α	22622	1	1233	MANDLPHSNSGDGQKPSSPDLF
		ŀ		1		TAPLLADPTVGFLPNDAEELFIF
				l	İ	LTEITEITIPCRVTDPQLVVTLHE
		l		ĺ		KKGDVALPVPYDHQRGFSGIFE
				!		DRSYICKTTIGDREVDSDAYYV
				l		YRLQAVPPVQLLTGWCVTAKA
		l		į.		PPDISAISALTAVKHGNCSSLTP
1		l				LLNPPGSDVIVCAEMDEQWGY
1		l		ŀ		VGAKSRQRWLFYAY\DSLRAR
İ		1				RVVAHVFG\*RTMATLGRLMSL
		l		Ì		LSPFDVVIWMTDGWPLYESRL
		1	1			KGKLHVISKRYTQRI\ERHNLNL
		l				RQHLARLGRKSLSFSKSVELHD
						KPSPQHREGVKDVMRFNEPMY
		1	ĺ			PLESRPRLYVCHLRNGAQASGR
		l	1			RCQLPMGCACHPVSQRNDRPN
		l			l	RRPERHATRKTTTRQRQQNSRQ
	1	1		l		HKQTQKKAKPNKRRATERGKG
	L	<u></u>	l		l	PQQKAERPKRHTHCFLKLQ

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nuclcotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
l	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22495	52863	A	22623	3	1386	HEASSPLAMASLTVKAYLLGK
22493	32803	^	22023	ľ	1360	EDAAREIRRFSFCCSPEPEAEAE
						AAAGPGPCERLLSRVAALFPAL
						RPGGFQAHYRDEDGDLVAFSS
l						
l					Į.	DEELTMAMSYVKDDIFRI*IKE
						KKECRRDHRPP\CAQEAPRNMV
						HPNVICDGCNGPVVGTRYKCS
l						VCPYY\DLCSVCEGKGLHRGHT
						KLAFP\SPFGHL\SEGF\SHSRWL\
					'	RKV\KHGHF\GWPGW\EMGPPG\
					1	NWEPTFLPRAG\EARPGP\TAES
ı						ASGPSEDPSVNFLK\NVGESVA\
			0			AAFSPL\NFEVDIDVEHGGKRSR
			0			LTPVSPESSSTEEKS\SSQPSSCC
			١.			SDPSKPGGNVEGATQSLAEQM
						RKIALESEGRPEEQMESDNCSG
				l	i	GDDDWTHLSSKEVDPSTGELQ
						SLOMPESEGPSSLDPSQGGNPQ
				ĺ	i	GWKEAALYPHLP\PEVEPAA*F
						ESLSOMV VPWGFS*LKGGWVH
						QGSCRPRNYDI\GAAL\DTIPVIQ
22496	52864	Α	22624	1032	1127	FYAKAEAGDRTVA*PHKPKMF
						LSIHHRVAKI
22497	52865	A	22625	121	561	VFPGPLGSLVDICGPSRVPFPWV
						VPAVLPVOFVORLPRFFRCPPV
					1	LVTSVIRSISTLIQALGSSPFARR
l					ĺ	YWGNLG*FLFLGVLRCFSSTGS
			-			PH*PMNSVNDSVSKHT/EVTPF
l			Ì			GN/PGNNGSYHLTDAYRRLARP
						SSPLT\PGHPPLYG
22498	52866	Α	22626	243	454	ATAGCPSCSATEGVVRNGKST
12450	12000	ľ	22020		[	AGHORYLCSPCRKTWOLOFTY
l					ľ	TASQP/GARIMGVGLNTVLRHL
	1		1		1	KNSGRSR
22499	52867	Α	22627	3	640	
22500		_	22628	236	410	
		-		116		
	52868 52869	A A				

NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
222502	52870	A	22630	276	1905	IFI YSSHQYGCPSCSATDGVV RNGKSTAGHQRYLCSHCRKTW QLQFTYTASQPAGQ*RTAIATVI CCNNCQRRHITNSRCHTDNCRR ADSASSHATSGSSRGCAGSGC CGASSTRRCLGPVCCFRGSKRS WRRQHEGHIKDCRKLVFHNFSFI HRMDNTAVPPVQLLTGWCYTA KAPPDISAISALTAVKEGNCSSL TPLLNPPGSDVICAEMDEQW GYYGAKSRQRWLFYAYDSLEK KAPPDISAISALTAVKEGNCSSL LSPFDVVIWMTDGWPLYSERL KGKLHVISKRYTQRIERHNLNL RQHLARLGRKSLSFSKSVEH KGKLHVISKRYTQRIERHNLNL RQHLARLGRADALDAPLAVDAQTP LSELLSHYGQAPCAVPVVGMEL ZEGLISHYGQAPCAVPVVGMEL GYVGIISKGMLLRALDREGGV RYPVIYFKRRRITDVCMPLSIG RVPVIYFKRRFRTDVCMPLSIG EKLKITPPERDSIRLDKPVML GGPLAEDRGFILHTPASNFASSI RISDNTYMTTSRDVLETLGTDK GYBDVLVALGYASWEKGQLEQ EILDNAWLTAPADLDILFKTPIA RYBCARKICYDLTMPAQD GTPDWWIIERLLKEWOPDEIIVG LPLNMDGTEQPLTARARKFAN RHGRFGVEVKLHDERLSTVEA RSGLIEEGGGYRALNKGKVDSA SAVIILESYFEGGY

NO: of peptide sequence hod in USSN (09540,217 location of first period sequence) in USSN (09540,217 location of preptide sequence) for peptide sequence period deletion, =possible nucleotide in peptide sequence period deletion, =possible nucleotide in peptide sequence period deletion, =possible nucleotide in period deletio	wn,
22503 52871 A 22631 399 2337 PEVQGLQQAGAAARPC APRVVCEGRLRGSGLQS ASV*ATDGVVRGKSTA YLCSHCRKTWQLQFTYY AGG*REKEEKEEKEEKE KRRREGEKEEKEEKE FKKKKKKKKKKKKKKKKKKKK KKKKKKQLSIVTMS WRNEARTTGACLPSPTY PIVSTTYTAFAAATLAK RASSKQVPLLVVLVWVY WFVKAGSVALGSSPGCV CKLLTGWCYTAKAPPDI LTAVKHGNCSSLTPLLNI VIVCAEMDEQWGYVGA RWLFYAYDSLRKTVVAH RTMATLGRLMSLLSPFD MTDGWPLYESRLKGKLI KARSKQQKDADTAPPQI TPGTYHAARDTRQASPR QHRQADGDKRQTRHAE RTHTGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTEVAARQTTEKDNEGS TQKTPSATATAHTPRTT VQARHTSAREENHTRQH RETTTAMTNTPHYHAE SQQQTTSTERPTIDSSTT. HTPRTVROERRMTQHEK STTTNRTTTTGATQAYQHTQI	
22503 52871 A 22631 399 2337 PEVQGLQQAGAAARRPC APRVVCEGRLRGSGLQS ASV*ATDGVVRNGKSTA YLCSHCRKTWQLQFTY1 AGQ*REKEEKEEKEEKEE FKKKKKKKKKKKKKKKKKKKKKKKK	ertion)
APRVVCEGRLRGSGLQS ASV*ATDGVVRNGKSTA YLCSHCRKTWQLQFTY1 AGQ*REKEEKEEVEEEEE KRREGEKEEKEEKEE KRREGEKEEKEEKEE FKKKKKKKKKKKKKKK KKKKKKKKKK	
APRVVCEGRLRGSGLQS ASV*ATDGVVRNGKSTA YLCSHCRKTWQLQFTY1 AGQ*REKEEKEEVEEEEE KRREGEKEEKEEKEE KRREGEKEEKEEKEE FKKKKKKKKKKKKKKK KKKKKKKKKK	VGLA
ASV*ATDGVVRNGKSTÄ YLCSHCRKTWQLQFTYT AGQ*REKEEKEVEEEEI KRRREGEKEEKEEKEEKEE EKKKKKKKKKKKKKKKK KKKKKKKK	
YLCSHCRKTWQLQFTYY AGQ*REKEEKEEVEEEEE KRRRGGEKEEKEEKEEKEE FKKKKKKKKKKKKKKKKKKKKKKK	
AGO*REKEKEKEVEEEEE  KRREGEKEKEKEKEKE KRKKKKKKKKKKKK KKKKKKKKKK	
KRREGGKEEKEKEEKEKE FKKKKKKKKKKKKKKKKKKKKKKK	
FKKKKKKKKKKKKKKK  KKKKKKKKKKKKKKKKKK  KKKK	
RKKKKK KQLSLVTVMS WRNEARTTGACLPSPTV PIVSTTVTAFAAATLAK RASSKQVPLLVVLVWVV WFVKAGSVALGSPGCV CKLLTGWCVTAKAPDDI LTAVKHGNCSSLTPLLNI VIVCAEMDEQWGYVGA RWLFYAYDSLRKTVVAI RTMATLGRLMSLSPFD MTDGWPLYESRLKGKLI KARSKQOKDADTAPPIQ TPCTYHAARDTRQASPR QHRQADDDKRQTRHAE RTHTGRKTHKETATQQ, QTRRTARTHRRTARHTA RTEVAARQTTEKDNEQSS TQKTPSATTAHTPRRTI VQARHTSARERNHTRQF RETTTNAHTHTPHYHAE SQQQTTSTERPTIDSSTT. HTTRTVROERRMTQHKFI STTTNRTTTTGAXYQHTQ	
WRNEARTGACLPSPTY PIVSTTVTAFAAATLAK RASSKQVPLLVVLVWVV WFVKAGSVALGSSPGCV CKLLTGWCYTAKAPPDI LTAVKIGNCSSLTPLLNI VIVCAEMBEQWGYVGA RWLFYAYDSLRKTVVAI RTMATLGRLMSLLSPFD MTDOWPLYESRLGKLI KARSKQOKDADTAPPIQ TPOTYHAARDTRAASPP QHRQADDDKRQTKHAE RTHITGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTEVAARQTEKDNEGS TQKTFSATATAHTPRTI VQARHTSARERNHTRQI RETTINAHTNIPHYHAE SQQQTISTERPTIDSSTT HTPRTVROERRMOHEKI STTTNRTTTTGAYQHTQI	
PIVSTTVTAFAAATLAKC RASSKQVPLLVVLVWV WFVKAGSVALGSSPGCV CKLLTGWCVTAKAPPDI LTAVKHGNCSSLTPLLNI VIVCAEMDEQWGYVGA RWLFYAYDSLRKTVVAI RTMATLGRLMSLLSPFD MTDGWPLYESRLKGKLL KARSKQKDADTAPPIQ TPQTVHAARDTRQASPR QHRQADGDKRQTRHAE. RTHTGRRKTIKETATOQ. QTRRTARTHRRTARHTA RTRVAARQTTEKDNEQS TQKTPSATTATHTPRITT VQARHTSARERNHTRQH RETTTNAHTHTPHYHAE SQQQTTSTERPTIDSSTT. HTTRTVRDERRMTQHKH	
RASSKOVPLLVVLVWVV WFVKAGSVALGSSPGCV CKLLTGWCVTAKAPPDI LTAVKHGNCSSLTPLLNI VIVCAEMDEQWGYVGA RWLFYAYDSLRKTVVAH RTMATLGRLMSLLSPFD MTDGWPLYESRLKGKLI KARSKOQKDADTAPPIQ TPOTYHAARDTROASPR QHRQADDDKRQTRHAE RTHTGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKDNEQS TQKTPSATTATHTPRTI VQARHTSARERNHTRQL RETITINAHTNIPHYHAE SQQQTTSTERPTIDSSTT. HTPRTVROERRMOHEKI STTTNRTTTTGAYQHTQI	
WFVKAGSVALGSSPGCV CKLLITGWCYTAKAPPDI LTAVKHGNCSSLTPLLNI VIVCAEMDEQWGYVGA RWLFYAYDSLRKTVVAI RTIMATLGRLMSLLSPFD MTDGWPLYESRLKGKLL KARSKQKDADTAPPIQ TPOTVHAARDTRQASPR QHRQADDKRQTRHAE. RTHTGRKTIKEFATAQQ, QTRATARTHRRTARHTA RTRVAARQTTEKDNEQS TQKTPSATATAHTPRRTT VQARHTSARERNHTRQH REITTIMAHTNFHYHAE SQQQTTSTERRTTDSSTT HITRTVRORRAMTKELSSTT	
CKLLTGWCVTAKAPPDI LTAVKHGNCSSLTPLLNI VIVCAEMBEQWGYVGA RWLFYAYDSLRKTVVAI RTMATLGRLMSLSSPED MTDGWPLYESRLKGKLI KARSKOQKDADTAPPIQ TPCTYHAARDTROASPR QHRQADDDKRQTRHAE RTHTGRKTHKETATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKDNEQSS TQKTPSATTAHTPRRTI VQARHTSARERNHTRQH RETTTNAHTMPHYHAE SQQQTTSTERPTIDSSTT. HTTRTVROERRMGHEKI STTTNRTTTTGAYQHTQI	
LTAVKHGNCSSLTPLINI VIVCAEMDEQWGYVOX  RWLFYAYDSLRKTVVAI RTMATLGRLMSLLSPFD MTDGWPLYESRLKGKLI KARSKQQKDADI'APPIQ TPQTVHAARDTRQASPR QHRQADDDRQTRHAE RTHTGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKOMEQS TQKTPSATATAHTPRRTI VQARHTSARERNHTRQH RETITINAHTHPHYHAE SQQQTTSTERPTDSSTT HTPRTVRDERRMTQHKI STTTNRTTTTGAYQHTQ	
VIVCAEMDEQWGYVGA RWLFYAYDSLRKTVVAH RTMATLGRLMSLLSPFD MTDGWPLYESRLKGKLL KARSKQKDADTAPPIQ TPQTYHAARDTRQASPR QHRQADDDKRQTRHAE. RTHITGRRKTIKEFATQQ. QTRRTARTHRRTARHTA RTRVAARQTTEKDNEQS TQKTPSATATAHTPRRTI VQARHTSARERNHTRQL RETITINAHTHTPHYHAE SQQQTTSTERPTTDSSTT HTRTVRDERRMTQHKL STTTNRTTTTGAYQHTQI	
RWLFYAYOSLRKTVVAH RTMATLGRLMSLLSPFD MTDGWPLYESRLKGKLI KARSKQQKDADTAPPIQ TPOTYHAARDTRQASPR QHRQADDDKRQTRHAE RTHTGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKDNEQS TQKTFSATTATHTPRTT VQARHTSARERNHTRQH RETTTMAHTNIPHYHAE SQQQTTSTERPTIDSSTT HTPRTVROERRMOHEKI STTTNRTTTTGAYQHTQ	
RTMATLGRLMSLLSPED MTDGWPLYESRLKGKLI KARSKQOKDADI'APPIQ TPOTVHAARDTRQASPR QHRQADGDKRQTRHAE. RTHTGRRKTIKETATQQ. QTRRTARTHRRTARHTA RTRVAARQTTEKOMEQS TQKTPSATATAHTPRRTI VQARHTSARESNHTRQI- RETITINAHTHPHYHAE SQQQTTSTERPTTDSSTT HTPRTVRDERRMTQHKI- STTTNRTTTTGAYQHTQI	
MTDGWPLYERLKGKLI KARSKQKDADTAPPIQ TPQTYHAARDTRQASPR QHRQADDDKRQTRHAE RTHITGRRKTIKEFATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKDNEQS TQKTPSATTAHTPRRTI VQARHTSARERNHTRQH RETITINAHTNPHYHAE SQQQTTSTERPTIDSSTT HTRRTVROERRMGHEKH STTTNRTTTTGAYQHTQ	
KARSKQKDADTAPPIQ TPOTYHAARDTRQASPR QHRQADGDKRQTRHAE RTHTGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKDMEQS TQKTPSATATAHTPRRTIT VQARHTSARERNHTRQH RETTINAHTNTPHYHAE SQQQTTSTERPTTDSSTT HTTPTVRDERRMTQHKI STTTNRTTTTGAYQHTQ	
TPOTVHAARDTROASRE QHRQADGDKRQTRHAE. RTHTGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKDMEQS TQKTPSATATAHTPRRTT VQARHTSARERNHTRQH RETITNAHTNPHYHAE SQQQTTSTERRTTDSSTT HTTPRTVRDERRMTQHKH STTTNRTTTGAYQHTQ	
QHRQADDKRQTRIAE RTHTGRRKTIKETATQQ, QTRATARTHRATARHTA RTRVAARQTTEKDNEQS TQKTPSATATAHTPRRTI VQARHTSARERNHTRQH RETITIAMHTNJYHYHAE SQQQTTSTERPTIDSSTT HTTRTVRDERRMTQHK	
RTHTGRRKTIKETATQQ QTRKTARTHRRTARHTA RTRVAARQTTEKDMEQS TQKTPSATATAHTPRRTI VQARHTSARENHTTRQI RETITINAHTNTPHYHAE SQQQTTSTERPTTDSSTT HTPRTVRDERRMTQHKI STTTNRTTTTGAYQHTQI	
QTRRTARHTARTARHTA RTRVAARQTTEKDNEQS TQKTPSATATAHTPRRTI VQARHTSARERNHTRQH RETITINAHTNIPHYHAE SQQQTTSTERPTIDSSTT. HTPRTVRDERRMIQHAY	
RTRVAARQTTEKDNEQS TQKTPSATATAHTPRRTI VQARHTSARERNHTQH RETTINAHTNTPHYHAE SQQQTTSTERPTIDSSTT HTPRTVRDERRMTQHKI STTTNRTTTIGAYQHTQI	
TQKTPSATATAHTPRRTI VQARHTSARERNHTRQI RETITINAHTINPHYHAE SQQQTTSTERPTTDSSTT HTPRTVRDERRMTQHKI STTTNRTTTTGAYQHTQI	
VQARHTSARERNHTRQH RETITINAHTNIPHYHAE SQQQTTSTERPTIDSSTT HIPRTVRDERRMTQHKI STTINRTTITGAYQHTQI	
RETTINAHTNIPHYHAE SQQQITSTERPITIOSITI HIPRTVROERRMIQHKI STITNRITITIGAYQHTQI	
HTPRTVRDERRMTQHKH STTTNRTTTTGAYQHTQI	•
HTPRTVRDERRMTQHKH STTTNRTTTTGAYQHTQI	INNNN
22504 52872 A 22632 103 313 ATAGCPSCSATEGVVRN	GKST
AGHORYLCSPCRKTWOL	
TASOP/GARIMGVGLNTV	
KNSGRSR	
22505 52873 A 22633 620 832	

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		Sequence		
22506	52874	Α	22634	453	1047	AADDRGHLRPTDGLLRNGKST
		l		1		AGHQRYLCSHCRKTWQLQFTY
		l				TASQPAGQ*RVTGTSERREQIIQ
		ĺ				RLRQQGSVQVNDLSALYGVST
		1				VTIRNDLAFLEKQGIAVRAYGG
		ı				ALICDSTTPSVEPSVEDKSALNT
		1				<b>AMKRSVAKAAVELIQPGHRVIL</b>
-						DSGTTTFEIARLMRKHTDVIAM
		1				TNGMNVANALLEAEGVELLMT
						AGICALLTGCCVTAKAPPDISAI
						SALTAVKHGNCSSLTPLLNPPG
l .			1			SDVPVCAEMDEQWGYVGAKS
		1				RQRWLFYAYDSLRKTVVAHVF
		1				GERTMATLGRLMSLLSPFDVVI
1		l				WMTDGWPLYESRLKGKLHVIS
		l				KRYTQRIERHNLNLRQHLARLG
		l				RKSLSFSKSVELHDKVIGHYLN
						GDFLQSTLREAPLIPHTREHP
22507	52875	Α	22635	1089	1686	TSPGYWRANSTDPDGRYCRTG
		ĺ				TSGGVCHALSSLRCPSCSATDG
1						VVRNGKSTAGHQRYLCSHCRK
		ı				TWQLQFTYTASQPAGQ*RFLSK
		l				QEKTLLINQLASEEQQKVTSRL
						SDALRNGRVWQLAIIYLTIQVA
						VYGLIFFLPTQVAALLGTKVGF
						TASVVTAIPWVAALFGTWLIPR
						YSDKTGERRNVAALTLLAAGIG
						IGLSGLLSPVMAIVALCVAAIGF
Į						IAVQPVFWTMPTQLLSGTALAA
					1	GIGFVNLFGAVGGFIAPILRVKA ETLFASDAAGLLTLAAVAVIDR
		l				RGTSLNRLRNTMIALLSNPFKE
						RLRKGEVOIGLWLSSTTAYMA
		1				EIAATSGYDWLLIDGEHAPNTI
						QDLYHQLQAVAPYASQPVIRPV
		ı				EGSKPLIKQVLDIGAQTLLIPMV
		1				DTAEQARQVVSATRYPPYAVH
		l				PVQLLTGWCVTAKAPPDISAIS
						ALTAVKHGNCSSLTPLLNPPGS
		1				DVIVCAEMDEQWGYVGAKSR
1		l				QRWLFYAYDSLRKTVVAHVFG
1		1				ERTMATLGRLMSLLSPFDVVIW
		1				MTDGWPLYESRLKGKLHVISK
						RYTORIERHNLNLROHTGTGW
						DGSRCRSQKSGGAA
22508	52876	A	22636	91	200	SGIIAFQYLLMTLKVLPVASVSI
						SCPS\CSATDGVVRNGKSTAGH
		ĺ				QRYLCSHCRKTWQLQFTYTAS
		ĺ				QPAGQ*RGGA

22509   52877   A   22637   2649   3270   GAPVASVSISC NOKSTAGHQI LQFTYTASQP KNKHMTITEF GEISPINITAM LADIEAEDLA WRYLNRILLS VSYQTPTREIN RENGGIOQRW VPGSFNDQFA WYQRVFIPK DAVTHYGKV DAVTHYGKV DAVTHYGKV DAVTHYGKV	PSCSATDGVVR RYLCSHCRKTWQ AGQ*RTIGRAML VAQRMRDAQQK ITSRLLLDLTYGV RKRRLLRITRDD SLFRRAGDNCAR KKLDGLWAFSLD WESALQESRAIA DADIRNYAGNV
22509   52877   A   22637   2649   3270   GAPVASVSISC NOKSTAGHQI LQFTYTASQP KNKHMTITEF GEISPPINTAM LADIEAEDLA WRYLNRILLIS VSYQTPTERIN RENGGIOQRW VPGSFNDQFA WYQRVFIPK DAVTHYGKV DAVTHYGKV DAVTHYGKV DAVTHYGKV	PSCSATDGVVR RYLCSHCRKTWQ AGQ*RTIGRAML VAQRMRDAQQK ITSRLLLDLTYGV RKRRLLRITRDD SLFRRAGDNCAR KKLDGLWAFSLD KKLDGLWAFSLD DADIRNYAGNV
22509 52877 A 22637 2649 3270 GAPVASVSISG NGKSTAGHQI LQFTYTASQP KNKHMITTEF GEISPDINTAM LADIEAEDLA WEYLMRLILS VSYQTPTREIR RENGGIORW VPGSFNDQFA WYQREVFIFK DAVTHYGKV DAVTHYGKV DAVTHYGKV DAVTHYGKV DAVTHYGKV	RYLCSHCRKTWQ AGQ*RTIGRAML VAQRMRDAQQK ITSRLLLDLTYGV BKLFRLRITRDD BKLFRAGDNCAR KKLDGLWAFSLD /WESALQESRAIA DADIRNYAGNV
NGKSTAGHQI LQFTYTASQP KNKHMTITEF GEISPDINTAM LADIEAEDLA WRYLNRILILS 'VSYQTPTREIN RENGGIDQRW VPGSFNDQFA WYQREVFIPK DAVTHYGKV	RYLCSHCRKTWQ AGQ*RTIGRAML VAQRMRDAQQK ITSRLLLDLTYGV RKRRLRITRDD RKRRAGDNCAR KKLDGLWAFSLD /WESALQESRAIA DADIRNYAGNV
NGKSTAGHQI LQFTYTASQP KNKHMTITEF GEISPDINTAM LADIEAEDLA WRYLNRILILS 'VSYQTPTREIN RENGGIDQRW VPGSFNDQFA WYQREVFIPK DAVTHYGKV	RYLCSHCRKTWQ AGQ*RTIGRAML VAQRMRDAQQK ITSRLLLDLTYGV BKLFRLRITRDD BKLFRAGDNCAR KKLDGLWAFSLD /WESALQESRAIA DADIRNYAGNV
LQFTYTASQP KNKHMTITEF GEISPDINTAM LADIEAEDLA WRYLMRILLIS VSYQTPTREIF RENCGIDQRW VPGSFNDQFA WYQREVFIPK DAVTHYGKV DAVTHYGKV	AGQ*RTIGRAML VAQRMRDAQQK ITSRLLLDLTYGV RKRRLRITRDD SLFRRAGDNCAR KKLDGLWAFSLD WESALQESRAIA DADIRNYAGNV
KNKHMTITEF GEISPDINTAM LADIEAEDLA WRYLNRILLIS VSYGTYREH RENGGIOQRW VPGSFNDQFA WYQREVFIPK DAVTHYGKV	VAQRMRDAQQK ITSRLLLDLTYGV RKRRLLRITRDD SLFRRAGDNCAR KLDGLWAFSLD /WESALQESRAIA DADIRNYAGNV
GEISPDINTAM LADIEAEDLA WRYLMRILLIS VSYQTPTREIN RENGGIOREW VPGSFNDQFA WYQREVFIFK DAVTHYGKV	ITSRLLLDLTYĞV RKRRLLRITRDD SLFRRAGDNCAR KKLDGLWAFSLD WESALQESRAIA DADIRNYAGNV
LADIEAEDLA WRYNRILLIS VSYQTPTREII RENCGIDQRW VPGSFNDQFA WYQREVFIPK DAVTHYGKV	RKRRLLRITRDD SLFRRAGDNCAR KKLDGLWAFSI.D WESALQESRAIA DADIRNYAGNV
WRYLNRILLS . VSYQTPTREIN . RENGGIOQEW VPGSFNDQFA WYQREVFIPK DAVTHYGKV	SLFRRAGDNCAR KKLDGLWAFSLD WESALQESRAIA DADIRNYAGNV
. VSYQTPTREIK RENGGIDQRW VPGSTDQFA WYQREVFIRK DAVTHYGKV	KLDGLWAFSLD WESALQESRAIA DADIRNYAGNV
RENCGIDQRW VPGSFNDQFA WYQREVFIPK DAVTHYGKV	/WESALQESRAIA DADIRNYAGNV
VPGSFNDQFA WYQREVFIFM DAVTHYGKV	DADIRNYAGNV
WYQREVFIPK DAVTHYGKV	
DAVTHYGKV	
22510 52878 A 22638 II 2470 MNADTDYSIA	EAAFNKGETAM
	IDTSKVNYGVT
	KPFVGVLSAGIN
	EFLENYLLTDEG
	GAVALKSYEEE
	MENAQKGEIMP
	AVRTAVINAAS
	DAQTRITNCCRI
	RKNPMDVIKKKH
	SVLGLLGLLVG
	GEYLFAITTLILS
	KAYAWRYVYPG
	FPLVCTIAIGFNN
	MPAKSSSLRSSQ
	AERTRYDSPGSY
	RGGSPFGGPGKY
	IAPSAQNQSPHG
	HALDWLDAGYT
	PYFKLAYVYDDS
	SIDNGTEGSAVR
	KNFSAYTDANYL
	SANDLTGITAKD
	QEFGKLDKCLS
	EKQVIFSSDVNN
	SLRREENVLIGIT
	GLMKNDIPYKLI
	PAKQTITLIEQN
	LAWHNRILELRD
	QFAHIIFMIAAM
	FSSGQHSVARKS
	CHEGFTISANPNS
SPRYPTIIEQAI	EQPRIVDFSIVRF
22511 52879 A 22639 I 852	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
22512	52880	A	22640	Ī	747	MILIDPRYTDTGAGREDEWIPIR
		1				PGTDAALVNGLAYVMITENLV
						DOAFLDKYCVGYDEKTLPASA
		1				PKNGHYKAYILATD/GVVRTAK
						STAGHORYLCSHCRKTWOLOF
						TYTASQPAGQ*RYTDTGAGRE
						DEWIPIRPGTDAALVNGLAYV
						MITENLVDQAFLDKYCVGYDE
	1					KTLPASAPKNGHYKAYILATDG
	1				1	WCVPPKAPPDISAISALTAVKH
						GNCSSLTPLLNPPGSDVIVCAE
1		1				MDEOWGYVGAKSRORWLFYA
						YDSLRKTVVAHVFGERTMATL
						GRLMSLLSPFDVVIWMTDGWP
						LYESRLKGKLHVISKRYTQRIE
						RHNLNLRQHLARLGRKSLSFSK
						SVELHDKVIGHYLNIKHYO
22513	52881	A	22641	1722	2463	NPRLAGSEGSCPSCSATDGVVR
						NGKSTAGHQRYLCSHCRKTWQ
						LQFTYTASQPAGQ*RKLTPGCL
		1				FVALKGERFDAHDFADQAKAG
		1				GAGALLVSRPLDIDLPQLIVKD
			1			TRLAFGELAAWVROOVPARVV
		1				ALTGSSGKTSVKEMTAAILSQC
				1	l l	GNTLYTAGNLNNDIGVPMTLL
		1				RLTPEYDYAVIELGANHQGEIA
						WTVTHPPRFVTKQWLCFLPDG
		ı				LLLFAEARFVIWQMQPAVMAV
		1			i	DVAAFAHLFNQQQYQRFTGRP
		1				PDSSKLPWKITCNGECVTFHW
	1	1			i	WLLGVAVRISRLLPVRVMTAL
1		1				EGNYNGTEGISALPFNGIILAHS
l	-	1			l	NESEWVTFRNNKNNEAFLDRV
	1	1				YIVKVPYCLRISEEIKIYEKLLN
		1				HIFECWQHFLNNPESLQCYIWA
ļ		1			l	ANADFREARQVIFKTFFTVGVE
ł		ı				EELGIRQTRTNDFLVTGDNLLRI
		1				FRFDVGNEDKVRQQFAVVRIH
	ŀ	1				REVFLVTFHGVNQRFSRHREEF
1		1				LFEFCVSVTANLTRCFFGIGFNL
	ŀ	1				RLTFFVIGDDFPTLTQNFRILIGV
1	1				i	IDGEFRLAHKAVAANHAIGLNA
1						QNGCRNDFVAHAAGSRALAAD
	1	1			1	GRSLEHAPTVETVVAGPLCESG
						DVFTQQEGGNVETRALPEVKA
					1	AVPPVQLLTGWCVTAKALPDIS
						AISALTAVKHGNCSSLTPLLNPP
1	1				1	GSDVIVCAEMDEQWGYVGAKS
	1				I	RQRWLFYAYDRLRKTVVAHRL
			-			

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22514	52882	Α	22642	90	1254	KGLCSHSEMVVPSYSATDGVV
		l				RNGKSTAGHQRYLCSHCRKTW
	İ	1				QLQFTYTASQPAGQ*RDIVLTK
		ı				LMMVWHSQSGYEKAYAHIQR
		1				WWYRAILLLTGWCVTAKALPD
		l				ISAISALTAVKHGNCSSLTPLLN
	ì	1				PPGSDVIVCAEMDEQWGYVGA
		l				KSRQRWLFYAYDRLRKTVVAH
	1	l				VFGERTMATLGRLMSLLSPFDV
	1	1				VIWMTDGWPLYESRLKGKLHV
						ISKRYTQRIERHNLNLSSTSILVL
		1				LNGATQFQDLGDLNKELEETY
		1				TDSKNHISCVGNDFGFNDIFSR
						YVEAVGREGDVLLGISTSGNSA
						NVIKAIAAAREKGMKVITLTGK
		1				DGGKMAGTADIEIRVPHFGYA
		l				DRIQEIHIKVIHILIQLIEKEMVN
		l				FTGLVQRGGGTGPHKDGWGIT
						FYEVAFEDVAVNFTQEEWALL
		l				DPWQKKLYRDVMLETYRNLAS
		l				VGDDDNIPSLREQVAHQRYFKT
		<u> </u>			1000	WHVEREYFSK
22515	52883	Α	22643	690	I233	RLQSLHYQGRGKPVPMRSYSLS FLFTISNSAGVFRCPSCSATDGV
	1					VRNGKSTAGHORYLCSHCRKT
	İ	l				WOLQFTYTASQPAGQ*RLLGP
		l				KNRPREQNWLLWCERGVMSRL
		l				OALYLREOGENNVKVDNFOTK
						VDSCTRPLAGQQFYPCIPDDAL
		l				VOPLEGGNEVFORREAGHITHE
			ļ			EDIVRHQHGGWRGANSRDDTP
1				l		GTVELFNGFNOOSAIAKTLRAF
		l				NATROHDHVIFAISHFDORRIR
İ						QQLHAARAGNRQVTIAGDAGS
		l				GDFDTAANQQIDSGDGFSLFTT
	1	1				RGEANQCRCAHIHSPFFLPFQT
			1			VRVFSAVPPVQLLTGWCVTAK
		1				APPDISAISALTAVKHGNCSSLT
		1		1		PLLNPPGSDVIVCAEMDEQWG
	1	1		1		YVGAKSRQRWLFYAYDSLRKT
						VVAHVFAVKPYWDTGGGFHC
	Ī	-		1		RWIYMQQEKELLQLSLQQGKY
	1		t			NOKRAAELLGLTYHOFRALLK

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22516	52884	A	22644	1314	1483	RIQRYCGKTGCPSCSATDGVVR  KOKSTGAHORYLCSHCRKTWQ  LQFTYTASQPAGQ*RMFNADG  SEVAQCGNGARCFARFVLKG  LUTNKRDIRVSTANGKMVLTVT  DDDLVRVNNGEPNEFPSAVPF  RANKAEKTYMRAAEQFILCGV  VSMGNPHCVIQVDDVDTAAVE  TLGPVLESHERPFERANIGFMQ  VVKREHIRLRVYERGAGETQA  COSGGACAAVAVGIQQGLLAEE  KYVELPGGRIDJAWKGEELQET  LTELDDRAVVDYLIKNPEFFIRN  RAAVEAIRVPHPVGGTVSLVE  WHMARARNHIHVLEENNALL  MEQAJANEGLFYBLLYLQRSLT  LAYAMPVTINKGNQIENQLDT  ETITISYEFILCOHLIANTEFSYL  ALPENYNRLCHPNSKNQTNNRF  KTLNSKAIGRILAAGGVYNGNI  EGRFTATEKLAVPPVQLLTGW  CVTAKALPPISAISALTAVKHG  NCSSLYTASOPAGO

NO:	of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
222517	52885	A	22645	1479	1793	TRYRSCPSCSATDGVVRNGKST AGHQRYLCSHCRKTWQLQFTY TASQPAGQ*RRNACQYRTPAV VASRQKRWHGOLGMIGSROV MLFISDSVGAKELDLIPRLEDK SIALLAMTGKFTSPLGLAAKAV LDISVEREACHMILAPTSSTVN TLMMGDALAMAVMQARGFNE EDFARSHPAGALGARLLNKVH HLMRRDDALPOVALTASVMDA MLELSRTGLGLVAVCDAQQAV QGVFTDGBLRRWLVGGGARLLNKVH PVNEAMTVGGTTLQSQSRAIDA KEILMKRKITAAPVVDENGKLT PVNEAMTVGGTTLQSQSRAIDA KEILMKRKITAAPVVDENGKLT VSHDRHRDAERQRVKSGLPRY AGAGLANRARNWYGSSWK VOYGOJILQGGPQNAEKSGQRVT VSHDRHRDAERQRVKSGLPRY AECSTYPSCPVALDLAFASHGI HMHAPGEALRMLVTAMDKAA DARTKLARLLATKGITHEIQIPD ISTKEKAQQFIGLNMEQIKPEKQ DFIKPVIPQAGKHNGVNMSVLR SLLTAGVLASGLLWSLNGITAT PAQASDDRYEVTQQRNPMSVLR SLLTAGVLASGLLWSLNGITAT PAQASDDRYEVTQQRNPMSVLR SLLTAGVLASGLLWSLNGITAT COYTAKAPPDISAISALTAVKHG CVTAKAPPDISAISALTAVKHG CVTAKAPPDISAISALTAVKHG CVSSLTPLINPPGSDVIVCAELD EQWGVYGAKSRQRWLFYAYA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22518	52886	Α	22646	807	1350	NRGCPSCSATDGVVRNGKSTA
				1		GHQRYLCSHCRKTWQLQFTYT
						ASQPAGQ*RLLNRLIEPTRGQV
						LIDGVDIAKISDAELREVRRKKI
						AMVFQSFALMPHMTVLDNTAF
						GMELAGINAEERREKALDALR
				l		QVGLENYAHSYPDELSGGMRQ
						RVGLARAINPDILLMDEAFSAL
		1				DPLIRTEMQDELVKLQAKHQR
						TIVFISHDLDEAMRIGDRIAIMQ
				ŀ		NGEVVQVGTPDEILNNPANDY
					l	VRTFFRGVDISQVFSAKDIARRT
						PNGLIRKTPGFGPRSALKLLQD
						EDREYGYVIERGNKFVGAVSID
						SLKTAAVPPVQLLTGWCVTAK
						APPDISAISALTAVKHGNCSSLT
						PLLNPPGSDVIVCAEMDEQWG
						YVGAKSRQRWLFYAYDSLRKT
						VVAHVFGERTMATLGRLMSLL
				i		SPFDVVIWMTDGWPLYESRLK
		l	ĺ			GKLHVISKRYTQRIERHNLNLR
						QHLARLGRKSLSFSKSVELHDK
22519	52887	A	22647		1270	VIGHYLNIKHYQ MTAATEOLNASRGLSTREMER
22319	32007	^	22047	l'	1270	OAELAKITTDYINSGGSEGDEK
						LONMIKAONDYYAAEDAKRA
						DWLAGAESAFADYGDSAMDM
						YGNVNEIASSALNGMTDMMV
						OFLTTGKANFADFAKNIISMIIK
						MIAOMVIFNTSLRNDGRSVSSG
						GAAVSLAAAAGSVATSGFNAS
		l	1			NSAPKVVTIREVERSVDVSGME
						ASSTSWITIRSFTAETSTFGPTM
						TASMTFLPTYCSIIASLLSSFKLA
						AATDGVVRNGKSTAGHQRYLC
i		İ	}			SHCRKTWQLQFTYTASQPALA
		1		l		RNPNRDITIYWGGREEQHLYDL
		1	1			CELEALSLKHPGLQVVPVVEQP
						EAGWRGRTGTVLTAVLQDHAT
		1		l		DGVVRNGKSTAGHQRYLCSHC
						RKTWQLQFTYTASQPGTHQKII
				ĺ		DMAMNGVGCRATAR\IMGVGL
	l	l				NTIFRHLKNSGRSR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22520	52888	IA	22648	381	945	GAPVASVSISCPSCSATDGVVR
22320	32000	^	22048	361	243	
						NGKSTAGHQRYLCSHCRKTWQ
				i		LQFTYTASQPAGQ*RTENQTPH
		ı				VLTHKWELNYENPWTQGGEHL
		l				TMGLVVGWRGLQEGDPARTN
		l				LSYYQANVGSGQHRENKGPIN
		1				LGNVGKSNGERGGKSLYRCKIF
			İ	l		LESSVTMKDRELPPVYDGVFEV
		1	l			LQWLLFLSAVPPVQLLTGWCV
		l		l		TAKAPPDISAISALTAVKHGNC
						SSLTPLLNPPGSDVIVCAEMDE
		1	1	l		QWGYVGAKSRQRWLFYAYDS
						LRKTVVAHVFGERTMATLGRL
		1				MSLLSPFDVVIWMTDGWPLYE
				l		SRLKGKLHVISKRYTQRIERHN
						LNLROHLARLGRKSLSFSKSVE
		ŀ		1		LHDKVIGHYLNIKHYQ
22521	52889	A	22649	354	1119	RRWRRGAHEAATGGTGRGPAE
22321	32007	l^	22047	1224	1117	EAAPGEPAAGRAAPKSGGKRL
						GGVLGRGGVGRRSGSPDRSGA
						HSAAPRRCRQLPGRCPSCSATD
			l			GVVRNGKSTAGHORYLCSHCR
						KTWQLQFTYTASQPAGQ*RSK
						GAIGEGPPDPRGIVEQNLGGDIS
	1		l		ľ	IPCYNGPRPRRSAQKTTLGPLN
				l		QPRIPSAQCAGAFPRVAAAGAG
				l		GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG
	1	1		l		RQRRRRWRRGEGEGDGDAGH
	1					MRPRPAGRAEARRRRRLRGNP
		1				PPGARLPNREGNGSGVSWAAE
						VSGDGRGARTEAARTRRPGG
				l		ADNYPGAVPPVQLLTGWCVTA
	1			1		KAPPDISAISALTAVKHGNCSSL
						TPLLNPPGSDVIVCAEMDEQW
					ľ	GYVGAKSRQRWLFYAYDSLRK
						TVVAHVFGERTMATLGRLMSL
				İ		LSPFDVVIWMTDGWPLYESRL
						KGKLHVISKRYTQRIERHNLNL
						ROHLARLGRKSLSFSKSVELHD
		l				KVIGHYLNIKHYQ
22522	52890	A	22650	23	658	VISRPGSSIQVRGLLLSRHGLAL
		Ľ			1	GSAPDLWKLPWVNLVQGMGQ
		1	l			OSOGSSCREGCSRPOCPSAAPC
1	1	1	l	I		GAAPAHFCSAATFWESEVQSIP
	1	1	1	I		R/CRATDGVVRNGKS/TAGHQR
				1		YLCSHCRKTWQ/LQFTYTASQP
	1	1		I		
		1		I		EVIRSSPTSAPRSDGTKTVPPLM
		1	1	1		GISP*SAHHRLATSDPSLCLSES
		1	1	I		LCSREPA A VPRASNQQPQGVK
		_				DVHHQFTARTSRYPSKC
22523	52891	A_	22651	926	1093	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22524	52892	A	22652	343	817	GGARSRQNPRCGPCMVVVPGQ
22324	32072	^	22032	545	017	SGRSVSADSSTEFREQGGS\AI\S
		l				FPKSINRAGPELFRFMENPFNSV
		l				*YRKNVRELTPAIMAVLPVEFR
				i		TRLAPONDTMSLIASAMKECSE
		ı				AKQAVLLNAPEHQKMKEVSEG
		l				IASLFRLMPEQVGPLMTMVTS
						MLGVI
22525	52893	A	22653	115	425	111111111111111111111111111111111111111
22526	52894	A	22654	747	1173	
22527	52895	Α	22655	I	2709	
22528	52896	Α	22656	1	2583	
22529	52897	Α	22657	636	1029	APSDQKRDFLVPHGADSAMAK
						SG*CESGNDLLSSSHMNQLR/A/
		l				ESIPEDVIAGASALVLTFIIKFVI
		l				AENPQWWQQFLKDHVSILAMN
						EDEAEALTGESDPLLASDKALD
						WVDLVLCTAGPIGLYMAGGGD
						VN
22530	52898	Α	22658	1611	3300	QSAWANRHRLVDCGRPGNGSD
						AGVAARSGRNDGTGRCGLCHS
						CSRYGDHHRQSDRIYRHYDAG
						RSPSGAVDYGTQRGNRFSRAV
						YPVGA/RRWR*GLPLV/QQFQPE
		1				NETSAAW\VVGINQPLVDIEAK
						VDDEFIERYGLSAGHSLVIEDD
				ł		VAEALYQDVQQKNLITHQSAG
						GTIGNTMHNYSVLADDRSVLL
				١.		GVMCSNIEIGSYAYRYLCNTSS
				l		RTDLNYLQGVDGPIGRCFTLIG
	l					ESGERTFAISPGHMNQLRAESIP
		l	ĺ			EDVIAGASALVLTSYLVRCKPG
		İ				EPMPEATMKAIEYAKKYNVPV
		l			1	VLTLGTKFVIAKNPQWWQQFL
					1	KDHVSILAMNEDEAEALTGESD
				1		PLLASDKALDWVDLVLCTAGPI
			l			GLYMAGFTEDEAKRKTQHPLL
		ĺ				PGAIAEFNQYEFSRAMRHKDC
						QNPLRVYSHIAPYMGGPEKCNI
			l	1		LGRSGWLLYGRGLDKQRLTQY
		ĺ		1		QSKLGAAMVIVAAWCVEDYQ
	1		l	1		VIRLAGSLTARATRLAHEAQLD
	1		1	1		VAPLGKIPHLRTPGLLVMDMDS
				1	1	TAIQIECIDEIAKLAGTGEMVAE
						VTRTGDARRTRFYRQPAQPCG
22531	52899	Α	22659	430	586	
22532	52900	Α	22660	1	131	

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1		ı		sequence		
		_				
22533	52901	Α	22661	ı	I282	MLPAHLDACVTTLQKIRWTLS
					ŀ	VVDPTLALRVPLPALSIRDDPPL
						RAYLPGRGALKVIRFDDMDFA
		l			ŀ	TIDGINRMLVDIDTNYLFLPRVT
		1				SSTIRCLSVPTRRTVPAVTASGR
						SVKYYPPDFDPSKIPKLKLPKD
						RQYVVRLMAPFNMRCKTCGEY
					1	IYKGKKF\NARKETVQNEVYLG
	i	l			l	\LPI\FRFYIK\CTRCLGRDSPFKT
1					1	DP\ENTDYTM/ESLGATRNFOG*
						EASWEEEEKRV\QK\EREDE/ES
						LNNPMKVLENRTKDSKLEMEV
						LENLQELKDLNQRQAHVDFEG\
						MLRQHRLSEEERRSQQQEEFEQ
					l	KTAALLEEARKRRLLEDSDSED
ł						EAAPSPLOPALRPNPTATLDEA
						PKPKRKVEVWEQSVGSLGSRPP
						LSRLVVVKKAKADPDCSNGQP
l						QAAPHPRSPAEQEGGQPYTPDA
						WRVLPEPTGCIPGQ
22534	52902	A	22662		412	WKVEFEFTGEFGQ
22535	52903	A	22663	298	532	MGSWSLPGRTLWKEKVRFLKS
22333	32303	n	22003	270	1552	KERTNPGSNFFSFPGSQHTREER
		1	l			SQECLSLPVLPSYLATRLSLPDS
		1	l		i	CPLKGRKAH*QRHSEI*LMGFE
		1			i	PGWPGKL*GWIELAFLLGCLHL
į.					1	TLPLPCPTPLPPKSTKSGCFSLSI
		ļ.				ALCFNKQFAVTK
22536	52904	A	22664	1	193	ALCFNKQFAVIK
22537	52904	В	22665	55	281	
22538	52905	A	22666	86	159	
22539	52907	A	22667	63	1036	VRGGLAAGRGRGSAGAAPVVV
22339	32907	A	22007	03	1030	AAMLGAWAVEGTAVALLRLL
		l				
i		l				LLLLPPAIRGPGLGVAGVAGAA
		1				GAGLPESVIWAVNAGGEAHVD
		1				VHGIHFRKCIPL*GRVGRALYY
		1				GMKLPILRSNPEDQSCYQTE\RS
		l				N/EETFGYESPIQREGGLTCWSL
		l				KFAE\VYFAQSQQKVFDVRLNG
		l	l	1		HVVVKDLDIFDRVGHSTAHDEI
		1	l		ĺ	IPMSIRKGKLSVQGEV\ST\FTQG
		1				KLLPLSFSKGYL*PIPKVCAL\YI
1	1	1	l	l	l	MAG\TVDDVPKASAFNPGLEK
		1	l	ĺ	i	KEEEEEEEYDEGSNLKKQTNK
		1				NRVQS\GPRTPNPYALDNSSLM
L		L_				FPILV\AFGVFIPTLFCLCRL
22540	52908	Α	22668	1570	1773	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
	L					L
22541	52909	A	22669	62	454	NKQLNKKRRRTMIALIQRVTRA
		1				S/VTVEGEVTGEIGAGLLV/LLG
						VEKDDDEQKAN/RDAEGQM/N
		1				LNVQQAGGSVLVVSQF/TLAAD
		1				TERGMRPSFSQGHHRIRAQAFY
	1	1		i .		DYFVES/CROOR*YAQGGLSAE
						FRLLNS
22542	52910	A	22670	1	1930	MAGKPEYDKTISTSIVLNALNA
	122710	ľ.		ľ		LGVSAEASGRNDLVVKTVEGD
		1				RKVSGSAYRETKDRGFHHGTL
		1	Į.	İ		LLNADLSRLANYLNPDKKKLA
				ł		AKGITSVRSRVTNLTELLPGITH
	1	1	ì	j		
İ			l	ĺ		EQVCEAITEAFFAHYGERVEAE
	l	1	ĺ			IISPNKTPDLPNFAETFARQSSW
						EWNFGQAPAFSHLLDERFTWG
		1		ŀ		GVELHFDVEKGHITRAQVFTDS
1		1				LNPAPLEALAGRLQGCLFSLYN
		1				SDLNTVIDNVLRIFPLLLSWISF
		1				WLLYSIVPTIRVPNRDAIVGAFV
	l	1			}	AALLFEAGKKGFALYITMFPSY
						QLIYGVLAVIPILFVWVYWTCV
	l	1	l			TVEGEVTGEIGAGLLVLLGVEK
				1		DDDEOKANRLCERVLGYRIFSD
l	l	1	1			AEG\KMNLNVQQAGGSVLVVS
İ	l	1				OFTLAADTERGMRPSFSKGVID
	1	1				DIGFNTEISQTRFHQASHLFKRK
	l	1				GIYRLHFARRRCQQIQRRQRRR
		1		l		RFMORTAVAITHOORMKOGIL
						RLWRNWLLWLLILLLRRNRLL
	l	1		ŀ		RLLILLLWRNRLLRLLILWLRR
	1	1		1		NRLLWLLILRLRRNWLLRLLIL
						LLRCNRLLRLLYRFNNWRKQR
						FVWTIIQQRFHWRKFKIIQRERI
						RPSRLFAGITLLLSLRKLTSQFS
		1	l			LCRSICIFCVYGHIMLILFAIALL
						LCLGELTCQFILLHRIDFIIVVLI
22543	52911	Α	22671	377	463	QTISQTTVQSN*NSGLRISPKTA
		1				QLHGD
22544	52912	Α	22672	129	615	KARQKSCQPVTTDTQCRHTSSS
1		1	1	1		IGKRLAFCQNLTQIHTESTMKP
1	1	1	ĺ	1		WILLLVMFISGVVMLLPVLGSF
1	1	1		1		WNKDPFLDMIRETEQCWVQPP
		1		1		YKYCEKRCTKIMTCVRPNHTC
1		1		1	1	CWTYCGNICLD/IRRAP*/HQC*
l						TPRFYWPITTVG*RMTIWSKKS
1		1		1		
22515	52012	١.	22672		412	AAYFLLGTPTWL
22545	52913	Α	22673	3	412	
22546	52914	Α	22674	2	390	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
22547	52915	Α	22675	180	507	CCGGRARSELCLLNLCIKGAD*I
		l		1		QAW*KFCQWPCERADGRGCRK
						AER\CPCPKGAYSL/YCRESVCC
						LPSFVNKNSSSLCVAHNWGRH
1		l				CYLWTAADVKQFISSHMFPVE
1						HL
22548	52916	Α	22676	3	227	GGGRARSELCLLNLCIKGAD*I
		1				QAW*KFCQWPCERADGRGS*F
		ı				DRGAEVCNQTDWQSIRSSQGG
						RDQRFSTERA
22549	52917	Α	22677	567	842	
22550	52918	Α	22678	1	1400	MGWGQGPDVQLMNGHERSPS
		1				HSPREAKKSGSPLPTDIHCFSPQ
1		ı			1	SQPAMVITGVPTAPAGASAKM
1						GLCATPITGACPLWLRGFRGWR
ł		1			İ	CEDRCEQGTYGNDCHQRCQCQ
1		1		1		NGATCDHVTGECRCPPGYTGA
						FCEDLCPPGKHGPQCEQRCPCQ
			ŀ			NGGVCHHVTGECSCPSGWMLS
		1			1	FPGWRPI*FSKSL*MQGTVCGQ
	1			l		PCPEGRFGKNCSQECQCHNGGT
		l		l		CDAATGQCHCSPGYTGERAAV
						PDVRK\CQDECPVGTYGVLCAE
l		l			1	TCQCVNGGKCYHVSGACLCEA
	İ	1		l		GFAGERCEARLCPEGLYGIKCD
						KRCPCHLENTHRGDDREKAEF
				i		GHLLIALEVQYTEISNCIEKSQE
		1				ASCYDIRMLKQPYGETNTEEEL
1	1			1		AVNYVYSIYASKEQIRYKHGRS
1	1					FANGHVKELMEGPSRRIAGSGG
l						EKYKRDEEKNTLLRGQNRGGR
1				1		GKNPKKTKAWVKIQGQNKIEIS
		L				GVGADRKRFFREK

SEQ ID SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
52919	A	22679		2210	MEÖKGEFKNDTLIYEFFHIVAK PHPLPIKSSTYTTFQSARVTRFF LDAEKECKCQEGQVEAALIAM GGTCQGTQRLSTAGESESATH MREPAVQCRENSOCYSVLAGR RRFFQGAASPLHPASVALTQLH VRWASTATIGQVSGFPGKIRVS PTINLGDSDYIFTASHWGLGHY GPAVHVHSQGGIDERGSNDSG RLYVTNPGGNPSWTPNIDRLA EDGVKLTQHISAASLCTPSRAA FLTGRYPVRSGMYSSIGYRVLQ WTGASGGIPTSETTFAKILKEK GYATGLIGKWHLGILNCESASD HCHHPHLHGFDHFYGMFPSLIK GDCARWELSEKRVNLEQKLNF LFQVLALVALTLVAGKLTHLISEY VGALIVHADCELMRNHTITEOP MCFQRTTPLIFLQEVASFLKRNE HGPFLLFVSLHVHIPLITMENF LGKILDFILDYGDNVSEMLDWFNS HGGFLLFUSCHVHIPLITMENF LGKILDFILDYGDNVSEMLDWFNS HGGSLENOLGNTQVGGWNGIY KGGKOMIGGWEGGIRVPGIFRW PGVLPAGRVIGEPTSLMDVFPT VVKLAGGEVPODRVIDGQUL VITAGREYBEPSLHVHIPLTMENF LGKSLHGLDSDHEFLMHVCER VLHAARWHQRDRGTMWKVHF VTPVPQPEGAGACYGKKVCPCF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22552	52920	A	22680	185	2091	GSTERDNMLHLHHSCVCFRSW
						LPAMLAVLLSLAPSASSDISASR
						PNILLLMA\DDHGVGDIGCYGN
	ŀ				i	NTMRTPNIDRLAEDGVKLTQHI
i	İ	l		İ		SAASLCTPSRAAFLTGRYPVRS
l						GMVSSIGYRVLQWTGASAGFT
					l	TN\ETTFAKILKEKGYATGLIGK
			İ			WHLGLNCESASDHCHHPLHHG
						FEHFYGMPFSLMGDCARWELS
						EKRVNLEQKLNFLFQVLALVA
						LTLVAGKLTHLIPVSWMPVIWS
						ALSAVLLLASSYFVGALIVHAD
		i				CFLMRNHTITEQPMCFQRTTPLI
						LQEVASFLKRNKHGPFLLFVSF
1		İ		ŀ		LHVHIPLITMENFLGKSLHGLY
		i				GDNVEEMDWMVGRILDTLDV
						EGLSNSTLIYFTSDHGGSLENQL
		1				GNTQYGGWNGIYKGGKGMGG
1		1				WEGGIRVPGIFRWPGVLPAGRV
1						IGEPTSLMDVFPTVVRLAGGEV
ļ						PQDRVIDGQDLLPLLLGTAQHS
ŀ						DHEFLMHYCERFL\HAARWHQ
l						RDRGTMWESPTL*PPCVSSQEG
						AGAC\YGRK\VCPCFGEK\VVHH
			ŀ			DP\PLL\FDLSRDPSETHILTPASE
		l			1	PVFY\QVMER\VQQAVWEHQR
		1				TLSPVPMQLDRLGNIWRPWMQ
ì		l				PCCGPFPLCWCLREDDPQINVC
		L				SEKLEPRFLNFVTQIETNQLAM
22553	52921	Α	22681	209	392	
22554	52922	Α	22682	1	513	
22555	52923	Α	22683	3	613	
22556	52924	A	22684	1	2115	

NO:	SEQ ID NO: of peptide sequence	Mei hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22557	52925	A	22685	53	1966	VGQPRGLRTLEKLTASVAGGA ANFFGDIERDLGPRDTSRGIRG GNLGDRGSSAEMAYASWQR WSPVEWARWAWHATAYTSSGDS SLLVLQGDSGKRSSDSEEAFET EYSTTPVKASPAPPPPPPEVIPE EVSTGPPEEPGWGSETVPVPD GPRSDSVEGSPFRTOAHSFSGVF DEDQPIASTGTDILDFDNIELVD TFQTLEPRASDAKNQGGKVNT FRKSTIGSVPISKSTLSRSLSLQA SOFDGASSSGNPEAVALAPDAY STESAKRVVPPASGGGRVQNSP PVGRKTLPLTTAPEAGEVTPSD SGGQEDSPAKGLSVRLEFDVSE SUSSWDNQGVENPPTKKIGKKP VAKMPLRFRMKKTPEKLDNT PASPPRSPAERDIPHKGTYTIF DIDKWDDPNFNPPSSTSKMQES PKLFQQSYNTDPOTCDESVDPF KTSSKTTSSFSKSPASFEPBASA MEANAVDGGGLNKPAKKKT PLKTMVEDVMSVCSLEDTFRY KKSPKRSPLSEDPSQDFTPAATP ETPPVISAVVHATDEEKLAVTN GKWTCMTVDLEADKQDYPPAATP ETPPVISAVVHATDEEKLAVTN QKWTCMTVDLEADKQDYPQPD
						YEIEYMEKIGSSLPQDDDAPKK QALYLMFDTSQESPVKSSTVR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
22558	52926	A	22686	662	2927	HLRRKRNLPRREGLSPALLEET
						PLEPAVGPKAACPLDSESAEGV
l	i					VPPASGGKRCLFTTAPEAGEVT
		1				PSDSGGQEDSPAKGLSVRLEFD
	ŀ	ŀ				YSEDKSSWDNQQENPPPTKKIG
	•					KKPVAKMPLRRPKMKKTPEKL
		l				DNTPASPPRSPAEPNDIPIAKGT
	l		İ			YTFDIDKWDDPNFNPFSSTSKM
		ŀ				OESPKLPOOSYNFDPDTCDESV
1		1				DPFKTSSKTPSSPSKSPASFEIPA
1	i	1				SAMEANGVDGDGLNKPAKKK
		l				KTPLKTDTFRVKKSPKRSPLSD
	i	l				PPSQDPTPAATPETPPVISAVVH
	ŀ	l			1	ATDEEKLAVTNOKWTCMTVD
		l			İ	LEADKODYPOPSDLSTFVNETK
	İ	l				FSSPTEELDYRNSYEIEYMEKIG
		l				SSLPODDDAPKKOALYLMFDT
i .	İ	l				SQESPVKSSPVRMSESPTPCSGS
		1	i			SFEETGSPLRNTAA\KNO\HPVP
		l			İ	TRDWAP*PRSSHLAGCQRKSSQ
		1				E/VELGGPWGLGHPFQEGDLKF
1		1				TASPRGSFASADALLSRLAHPV
		1				FSL/CGALDYL\EPNFRQKRNPP
		Ì				LFAQKLQREA\VHPTDVSI\SKT
1			l			ALYSRIRTTEVEKPAGLLFQQP
ł					i	DL\DSALQIARAEII\TKEREVSE
		l			İ	WKDKYEESR\REVMEMRKI\LA
					i	EYEKTIAQMIEDEQREKSVS\HQ
						TVQQLVLEKEQA\LADLNSVEK
				1		SLADLFRRYEKMKEVLESFRK
		l		ļ		NEEVLKRCAQEYLSRVKKEEQ
						RYQA\LKVHA\EEKLDRANAE\I
22559	52927	Α	22687	374	1268	AETNMVWYDWMRPSHAQLHS
						DYMQPLTEAKAKSKNKVRGV
		1		1		QQLIQRLRLIKSPAEIERMQIAG
				•		KLTSQAFIETMFTSKAPVEEAFL
		i				YAKFEFECRARGADILAYPPVV
		l				AGGNRSNTLHYVKNNQLIKDG
		i				EMVLLDGGCESSCYVSDITRTW
						PVNGRFTAPQAELYEAVLEIQR
	l		1		l	DCLALCFPGTSLENIYSMMLTLI
						GQKLKDLGIMKNTQGK*CL/RR
			İ			LLE/QYCPHHVGHYLGMDVHD
1		1	İ			TPDMPRSLPLQPGMVITIEPGIYI
			1			PEDDKDAPEKFRGLGVRIEDDV
	1	1				VVTQDSPLILSADCP
					L	

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22560	52928	IA	22688	329	1389	RIWKTOMMRMKKMMMMMM
22560	52928	A	22088	329	1389	
				i	1	MMMRTTLFFQQESQEAIFFPFD
		İ		1		LFPMCPFGCQCYSRVVHCSDLG
						LTSVPTNIPFDTRMLDLQNNKI
		1				KEIKENDFKGLTSLYGLILNNN
1						KLTKIHPKAFLTTKKLRRLYLS
						HNQLSEIPLNLPKSLAELRIHEN
					}	KVKKIQKDTFKGMNALHVLEM
i						SANPL**/SMGIEPGAF*RG*RCS
						/YIRIAEAKLTSV\PKGL\PPTLLE
					İ	LHLDYNKISTVELEDFKRYKEL
					'	QRLGLGNNKITDIENGSLANIPR
				ŀ		VR\EIHLGNIKLKKIPSGLPELKY
						LQIIFLHSNSIARVGVNDFCPTV
i						PKMKKSLYSAISLFNNPVKYWE
						MQPATFRCVLSRMSVQLGNFG
22561	52929	Α	22689	1	4608	
22562	52930	Α	22690	1	465	
22563	52931	Α	22691	1	1542	
22564	52932	Α	22692	1	4226	EAVLCSSRGRPDSSMPDCCRNA
		1				ALEARSCQSMTSLFSNTVSPTQ
		1			<b>!</b>	DGTSSLPRRQSSFAKPPLRALY
		l	l			DLLIAPMEGGVVESTEPLGTSR
		ł				SGFASGTALFWLGDNGKAASP
1		l				PGVSVSSLENEDDNTQCNRLEQ
						GYREAAVPDWTODLALCLLPT
		l				LOSLKARTAPHLCVPSSCYKA
		1	1			WYTLAVSKVSGMVEGKAWIIL
		1				LLLAASTHALLQRPQEGDITPL
		ı				AGSRAAAVSCOARPFSTFSGLM
		ı				HSSGPVGRHROLILVLE
22565	52933	A	22693	2	4197	ALDFPGRRFRPTASFIWASVLFE
22303	32,755	ľ`	22075	ľ	4157	TIRHEAEVSTDYKLSLFDLQTSS
		l				YQALQRVLVSLGHHDEALAVA
	1	ı	1			ERGRTRAFADLLVEROTGOOD
		ı				SDPYSPVTIDOILEMVNGORGL
		ı				VLYYSLAAGYLYSWLLAPGAG
		ı				
		l	1		1	IVKFHEHYLGENTVENSSDFQA
		l		1		SSSVTLPTATGSALEQHIASVRE
		I	1	l		ALGVESHYSRACASSETESEAG
		ı				DIMDQQFEEMNNKLNSVTDPT
		1		l		GFLRMVRRNNLFNRSCQSMTS
		L				LFSNTVSPTQDGTSS
22566	52934	Α	22694	1	618	
22567	52935	Α	22695	1	3528	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
22568	52936	A	22696	85	1193	RPIEGOLGTCHPGSEPRTIRWEA PGKGR LR VTEPPTSDIAGRTGA AASFPAPLHLPREPRAGVDGTD AERILYPRRAAPAASGVOSRPA GEA VRADEAGAAGLGGFQSER NRSWGGESLGSRQPOKMPAFN RLFPLASL VLIV*VSVCEPPWG VPLENGGRARANPMKLRCISC MKREEVEATTVVEWFYRPEGG LQWNGSKDLQDVSITVLNVTL NDSGLYTCNVSREFEFEAHRP VKTTRLIPLR VTEEAGEDFTSV VSEIMMYILLVELTL WLLIEMIY CYRQVSKAEEAAQENAFGNSL EGGSQGLQNNSTQDGLVTGQQ UVGSHRAREAGNENTDEGLYTGQD UVGSHRAREAMPDEFEEPPSAK
22569	52937	A	22697	277	527	VRVSWEYFCAPGPAGSPAGPG RGEEKQRLGLLIRDARSPAPPG PPVGSVRG/PKPAPAGRVRAKG RRA*GAACSLAAGAWVEPGI

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	· ·			sequence		· · · · · · · · · · · · · · · · · · ·
22570	52938	A	22698	226	2268	SVKNYTKCHVRNEQICNKLTSC
	1		l		1	KSCSLNLNCQWDQRQQECQAL
		ı	ŀ			PAHLCGEGWSHIGDACLRVNSS
		1	ĺ		1	RENYDNAKLYCYNLSGNLASL
		l			l	TTSKEVEFVLDEIQKYTQQKVS
						PWVGLRKINISYWGWEDMSPF
		l				TNTTLOWLPGEPNDSGFCAYLE
				ŀ		RAAVAGLKANPCTSMANGLVC
		l				EKPVVSPNONARPCKKPCSLRT
		l	ļ			SCSNCTSNGMECMWCSSTKRC
		l	ŀ			VDSNAYIISFPYGQCLEWQTAT
		l		l		CSPONCSGLRTCGOCLEOPGCG
		l	ŀ			WCNDPSNTGRGHCIEGSSRGPM
						KLIGMHHNEMVLDTNLCPKEK
					Į.	NYEWSFIOCPACOCNGHSTCIN
		1		İ		NNVCEOCKNLTTGKOCODCMP
		l				GYYGDPTNGGOCTACTCSGHA
		1	ŀ			NICHLHTGKCFCTTKGIKGDOC
					ŀ	OLCDSENRYVGNPLRGTCYYK
					ĺ	T\LIDYQFTF\SLLQEDDRHHTAI
		1			ļ	YFIVNPKQSNKNLHISINASNNF
					ŀ	NLNITWPC*STAGTISGEETYIV
						SKNNILEYRDSFSYEKFNFRSNP
					l	NITFYVYVSNFSWPIKIQIAFSQ
					1	HNTIMDLVOFFVTFFSCFLSLLL
		1				
		1			İ	EAAVVWKIKQTCWASRRREQL
		1				LRERQQMASRPFASVDVALEV
						GAEQTEFLRGPLEGAPKPIAIEP
						CAGNRAAVLTVFLCLPRGSSGA
						PPPGQSGLAIASALIDISQQKAS
00.581	50000	_	22522	2.52	***	DSKDKTSGVRNRKHLSTRQGT
22571	52939	В	22699	252	360	
22572	52940	A	22700	1	84	
22573	52941	A	22701	1	404	1000 to 00 000 000 000 000 000 000 000 00
22574	52942	A	22702	86	292	NPEMRCREFCLPSCRPELALE*
			1	1		G*SVRDVA*AQMCSVVWAGRP
						LGFAA*S*GSGHSCGRCSIWGE
		<u>_</u>		L		CWG
22575	52943	Α	22703	82	379	HTLQPFISMGSLTKMFKVISEGP
						MD*RPHPKVRPMT*HSTPAPAN
						FMGHFFWKIMICTDRTEVATLV
			1			AGV VSGASVGGKSPPYRKWNS
						PEPDVKGPRVL
22576	52944	Α	22704	2	382	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22577	52945	A	22705	1	479	HMPSGETAPLRIVDTGTGTGTV
						\AYAPTE\LGLHEMHIKYMASHI
						LESPLRFYVTTPTVPSVSAYGPG
						LVYGVAN/NTATFTIVTEDAGE
		1	ł			GGLDLAIEGPSKAEEPRPRLSRA
		1				PGVDGAGASPGLGEDTGPQAP
						GLQVLTERIPGPSAPTPEFLSVG VSAVC
22578	52946	A	22706	1	590	VSAVC
22579		A	22707	171	406	
	52947			269	2477	
22580	52948	A	22708		=	
22581	52949	В	22709	167	7472	
22582	52950	Α	22710	1	520	
22583	52951	Α	22711	2218	2472	
22584	52952	A	22712	722	1555	SRNWYEALNNTWGREDICVVD
ŀ						GWQSFRRQDYAFLQERGSGGD
		l				LWSSTLAQAMTLVESRHPRHQ
ļ.		1			- 50	ALSTQLLDAIMPYCGNTLRLGV
						TGTPGAGKSTFLEAFGMLLIRE
ŀ						GLKVAVIAVDPSSPVTGGSILG
		1				DKTRMNDLARAEAAFIRPVPSS
			ĺ			GHLGGASQRARELMLLCEAAG
	i	1				YDVVIVETVGVGO\SETEVARM
			i			VDCFISLOIAGGADDLOGIKKG
	1	1				LMEVADLIVINKDDGDNHTNV
	1					AIARHM/YESALHILRRPPMHR
				1		VPRNPKVNLLPGTAMANLLS
22585	52953	В	22713	381	425	
22586	52954	A	22714	1	1185	
22587	52955	A	22715	1	3009	
22588	52956	A	22716	22	527	YEGIVMKPLIIVNPADCIGC\RT
		1				CE\VACVVAHPSEOELNADVFL
						PRLKVORLDSISAPVMCHOCEN
1	1	ı	1	I	1	APCVGACPVGALTMGEQVVQT
l	1	1	1	1	1	NSARCIGCOSCVSACPFGMITIO
l	ĺ					SLPGDTROOIVKCDLCEOR*EG
		l				PACVESCPTQALQLLIERELRRV
l		l		I		ROORIVVTGENPL
22589	52957	A	22717	187	325	INQUITY TO ENTE
22309	36731	<u></u>	22/1/	107	323	

SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequenee	codon for last amino acid of peptide scquence	Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \*=possible nucleotide insertion)
52958	A	22718		2240	IMRSDPDTSVNEIETKLSALLGS ETTGEILFDLLCANGPEWNRFV TLEMKYGRIMLDTAKIIDEODV PTHILSKLTFTLRNHPEGVVMK NFEVLOPLOSULSGLPLWYSER ILQQINQLTHYEPVIGIMGKTGA GKSSLCNALFAGESPYSDVAA GKSSLCNALFAGESPYSDVAA GKSSLCNALFAGESPYSDVAA CTRDELRERLQIGEHFMTIVDLP GVGESGVRDTEYAALYREQLP GVGESGVRDTEYAALYREQLP GVGESGVRDTEYAALYREQLP YROUGEAYRHKMLFVISQSDK AEPTSGGNILSTEQKQNISSKIC LHELFQPWHPCAVSVSLHPV VALLQOFRTDDERTHRYHYDS QHRLVDYTRTQYEEPLVESRYL YPDLGRRYAKRVWRRERDLTG WMSLSKKPQVTWYGWDGDAL TIQNDRSKIGNTYQGGSFTPLIR VETATGELAKTQRRSLADALQ QSGGEDGGSVVPPVLVQMLD QUELSGLGGSVVPVLQQUL QLIRLPGQQYDEESGLYVNRH YYPDPLQGRYTTQDPGLKGGW VARQMNAIASKLPEGFHIGINFS ASHISPTFVDECLNFUSSHTQL SHILDFTFVDECLNFUTSSHTQL SHILDFTFVDECLNFUTSSHTQL SHILDFTFVDECLNFUTSSHTQL SHILDFTFVDECLNFUTSSHTQR SHILDFTFVDECLNFUTSSHTQR SHILDFTFVDECLNFUTSSHTQR SHILDFTFVDECLNFUTSSHTQR SHILDFTFVDECTHTQL SHILDFTFVDECTHTQL SHILDFTTPTTTPTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22591	52959	Α	22719	1	2301	MALTSGVHTGEKPYKCAQCEE
						AFSRKSELIIHQIIHTGEKPYECT
						ECGKTFSRKSQLIIHQRTHTGEK
	1					PYKCTKCGKSFCQQSHLIGHQR
		t				IHTGEQPYLGLSVAERPKEYED
						DRALKMAAPCGNSSREQPCCFL
		1				NKASEDAERAKPPS*EPGRLRT
			0.0			P*VAQPH*VPHFMATWVRKHP
	1	l			ŀ	VHDRQFRLHGQIRELNWDVVR
		l				PEQFTGVATNCSGITYPQGGWL
		l				CPAELTRNVLELAQQQGLQIYY
	1		İ			QYQLQNLSRKDDCWLLNFAGD
						QQATHSVVVLANGHQISRFSQT
		1	1			STLPVYSVAGQVSHIPTTPELAE
						LKQVLCYDGYLTPQNPANQHH
1		1				CIGASYHRGSEDTAYSEDDQQQ
		l				NRQRLIDCFPQAQWAKEVDVS
			İ	l		DKEARCGVRCATRDHLPMVGN
				l		VPDYEATLVEYASLAEQKDEA
		l		l		VSAPVFDDLFMFAALGSRGLCS
				l		APLCAEILAAQMSDEPIPMDAS
			1			TLAALNPNRLWVRKLLKGKAV
22592	52960	A	22720	1021	1734	KAG
22593	52961	Ā	22721	1247	2412	RRSHHAANLDIRIRMPQTRQVT
	12201	ľ		1.2		LPDPELHPDSTLSMWPD/NRIAR
ļ		1				DAHYLYRYDRHGRLTEKTDLIP
						EGVIRTDDERTHRYHYDSOHRL
		l				VHYTRTQYEEPLVESRYLYDPL
ĺ		l				GRRVAKRVWRRERDLTGWMS
İ	1					LSRKPOVTWYGWDGDRLTTIQ
						NDRTRIQTIYOPGSFTPLIRVET
1	1					ATGELAKTORRSLADALQQSG
						GEDGGSVVFPPVLVOMLDRLE
	ļ.			1		SEILADRVSEESRRWLASCGLT
	1					VEQMQNQMDPVYTPARKIHLY
						HCDHRGLPLALISKEGTTEWCA
						EYDEWGNLLNEENPHQLQQLI
					ĺ	RLPGQQYDEESGLYYNRHRYY
	1	1	l			DPLQGRYITQDPIGLKGGWNFY
			l	1		OYPLNPVTNTDPLGLEVFPRPFP
		1				LPIPWPKSPAQQQQWYPTFCC
22594	52962	Α	22722	1	4767	
-2007	1-2-02	۲.	1	1-	1	L

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22595	52963	Α	22723	668	1922	YPHADSPECWLSSSSERGKSSV
						TADLSRPRRLRPIPTCHGKPIAC
			İ			TTTKAKNAFRPIAGASATGRLA
						NSPIIILPNAAIRQEFGITLDPEEV
1						FTFQGVKLYEYRYFPDCVRSRN
1					}	EAGILVEQILSTLFSVGEHFSLK
						RAPFFKVPSLRSFYAFGGLTTG
						VVRRNGNIIGVNTPARKIHLYH
						CDHRGLPLALISKEGTTEWCAE
						YDEWGNLLNEENPHQLQQLIR
		l				LPGQQYDEESGLYYNRHRYYD
						PLQGRYITQDPIGLK\GGWNFY
	l					QYPLNPVQYIDSMGLASKYGH
	i					LNNGGYGARPNKPPTPDPSKLP
						DIAKQLRLPYPIDQASSAHNVF
			ŀ		İ	KTFFRALSPYDYTLYCRKWVK
1					Į.	PNLTCTPQDDSQYPGMDTKTA
						SDYLPQTN/WPVPSSIVSCTTGV
						VP*VVCQWVFAWPPDISHRRTG
22596	52964	Α	22724	25	510	RITDFGVKKIIRVGSCGAVLPHV
						KLRDV/VI/GMGAC/TDSKVNRI
						RFKGQAFGGIADFAMGRTAVD
						PGKALGINARVGNLVSADLFYS
	ļ					PDGEMFDVMEKYGILGVEMEA
	1					AGIYGVAAEFGAKALTICTVSD
	1					HIRTHEQTTAAERQTTFNDMIKI
						ALESVLLGDKE
22597	52965	A	22725	415	759	HRRRIVIDDTGIFRPRNTAQQLR
						QRPIAMSASGVVQIILEGHWRT
				į		HSLRNGFHRGFRL*SSPKVGMQ
						HGSRQVDHRTQRALRFFRQPGI
						DFLRPVFTPRRQRLAATNHFSR
	<u> </u>					FVEH

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22598	52966	A	22726	sequence 2	2329	GTLPGCGAMHTCPSGPATHLV RRRIVRLPACVVTRANVPTPSR PVNATIGAFHSG*MASQGRRLS AQYPWSPPQYAVARPVPTVPCR LVPTGTSKAFKISVMOFOMGIP SCSISAKELITDFGVKHIRVGSG GAVLPHVKLRDVVIGMGACTD SKVTRIRFKDHAFAAIADFDMV RNAVDAAKALGIDARVGNLFS SADLFYSPDGEMFDVMEKYGIL GVEMEAAGIYGVAAEFGAKAI TICTVSDHIRTHEQTTAABSQT FINDMIKIALESVLLGDKEEDRV RFGKARATRYALLRPYRGIERII VRWYDDTGKAHKFADIRLCWI QGSCLDVAKAREVIRGSQPLPA WCENWAQGVDARLTKEAQSA LRPVINLTGTVLHTNLGRALQA EAAVEAVAQAMRSPVTLEYDL DAGGGHRDRALAQLLCRITG AEDACIVNNNAAVLLMLAAT ASGKEVVVSRGELVEIGGAFRI PDVMRQAGCTLHEVGTTNRTH ANDYRQADFQEDDFVMAFTGM INGPYQNTRALHGGHRRPDFLP
			20505		200	LLGAFHGAIDILAPGGLHGYSL PSSFFHALALSAAFPGKRCKLS VHLPFWGLKDGGPLLTAPLGG ALVGIVGGVGGSDLTFPFHTHS LNVRFVIRRISHRHHHFYRIAKL LMNGAQGSNMLIAAVAGVPHC NGHFPFTVADRFAIVLSEGSIRV
22599 22600	52967 52968	A	22727	876	702 1318	RCHLANASPAAGSHAPSPDSLS KIR/QEVG/PPLP/PLLAPLIATPP RTSQPLSPLISSSSPSSPASPVGQ VSPFRETPVPPAMSPWPEDPRR ASPPDPSPSPSAASASERVVPSP LQFCAATPKHALPVPGRLPPCA SGHAMPCWRAKIK

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		•
22.601	10000	_	22729	485	1455	PVVORARNASSCOPFSTSVWAI
22601	52969	A	22129	483	1455	
		l			1	PSISATSVPMCGATHSILSPKKS
						TVSDRIGSMQISFFPLSRSAEK*
					1	ESPCSSDAFHAIFSVLSGLALHS
					i	TTTSLCSSTSGQLVCC**TSLPPT
					1	TYGMIACAAPVE*SPRCPV*PPA
						SAI*RCNNVEALCSTPFERQP*E
						PAK\SMPDHKPHECVYVVGE*D
						PAPHPNSREQIHPAHECLAGFP
		l			İ	VTQESLFAPLEIVPAYCCEPGRE
		l				PRSARSFPFAPRELT*VQLELGT
						GL*RETLDLVVSAGWA*ARRIS
		l				GA VTATPETGTLCQLDSGAECH
		l				YRETGVELP*RSPRPVGHGSSR
						HSRRWRT
22602	52970	A	22730	814	5001	PENSSHRPKPIIPMIASTLINANQ
						NSVSPYKRTFTRLTALMIIKKA
		1				AAQIHVGTSGNQYWRKLDFNS
		l				PGGSSPVENSDCSTNSRLSFSPE
	l	l				NILIQNQDIVREAAVQGDGQKQ
		l				RQPQATDLDSSGTHGSEMLPAT
		l				EVTVSGGFSVEETSCGDTGRSG
		l				GEALAVANDSTSTPQNANGLW
		l				KLKSTTPGGALPECFGTTDTTFS
		l				SAFCRKHGETQDTSQSSLPGTL
		l				HCYTGIREGGDDTEVESEAFSC
						SEGSEQQDAPD
22603	52971	Α	22731	34	102	
22604	52972	Α	22732	584	893	QKLLPNRMGHPVFVKSSQDPV
						LLSRIAWKDASTVTWVIEQDPV
		l				LRRRRRRTRRRRTRMRRRRGR*
		l				*/SKRRRMR/GRRGRR*CRRRRR
						RSSRSRPTRRQRNRNQV
22605	52973	Α	22733	2	269	IGKLYWFMPPSANLALRELGIV
			l			LFLSVVGLKSGLSWIGYGALIT
1		İ				AVPLITVGILARMLAKMNYLT
	l	l		,		MCGMLAGSMTDPPA*PRFQTN
1		l				DREEQHDPQLPQREVGA WRHK
		l				PVQLAD
			·		l	L`

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /-possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22606	52974	A	22734	3	958	RCYSLLSRAPHPGIYISIAERV/*I
22000	32914	^	22734	ľ	750	RLNSANFTLNSLLPAEKKLAEE
		1				FAVSRMTIRKAIDLLVAWGLV
						VRRHGSGTYLVRKDVLHQTAS
						LTGLGEVLKRQGKTVTSQVLIF
						EIMPAPPAIASQLRIQINEQIYFS
						RRVRFV\EGIPLMLEDSYMPVK
		1				LFRNLSLQHLEGSKFEYIEQECG
		1				ILIGGNYESLTPVLADRLLARQ
		1				MKVAEHTPLLRITSLSYSESGEF
						LNYSVMFRNATQIQGWRSHFR
					İ	R\LHPKKSLTVPPERAPPVFGGL
						IMRKNHSQRIDSGVRERRRTAG
		1		1		WMKIISKRORRRIGHGTCOHPA
		1				HGOVIHFG
22607	52975	Α	22735	2	285	GDEMTTSTLOKAIDLVTKATEE
		1				DKAKNYEEALRLYQHPVEYFL
					į.	HAVKYEAHSDKAKESIRAKCV
						QYLDRAEKLKDYLRSKEKHGK
						LCLKDRVT
22608	52976	A	22736	545	791	LCLRURVI
22609	52977	A	22737	249	1574	NHKAIDLVTKATEEDKAKNYE
22009	32711	^	22/3/	249	1374	EALRLYOHAVEYFLHAIKYEA
						HSDKAKESIRAKCVOYLYRAE
		1			ļ	KLKDYLRSKEKHGKKPVKENQ
		l				SEGKGSDSDSEGDNPEKKKLOE
					1	
						QLMGAVVMEKP\NIRWNDVAG
						LEGAKEALKEAVILPIKFPHLFT
	1					GKRTPWRGI\LLFRPPGT\GNSN
		l				RPKAGAPKATNSTFFSVSS\SDV
		ĺ	ŀ	l	l	KSKRLGETEKLV*NLF\ELARQ
				ł	1	HK\PSIIFIDEVDSLCGSRNENES\
i			İ		ŀ	QAARRIQNGSSLVQIAGGLGIN
ŀ				i		NDGDSGFFGATNIP\WVLD\SAI
	İ					R\RKV*KTKFYIPLPEKACPRPD
		i		l		VSRLHLGSTPHN\LRVATFHKL
						ARKTEGYSGADISIIVRDSLMQP
1		1	l	l		VRKVQSATHFKKVCGPSRTNPS
1		l		I		MMIDDLLTPCSPGDPRAMEMT
I		l	l	I	l	WMDVPGDKLLEPVVCMSDML
l		l	l	I		RSLATTRPTVNADDLL\KVKKF
						SEDFGQES
22610	52978	A	22738	52	567	
22611	52979	В	22739	1	1293	
22612	52980	Α	22740	443	1027	
22613	52981	A	22741	2077	2430	
22614	52982	Α	22742	1	504	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
					<u> </u>	
22615	52983	A	22743	3244	3947	RNEITPFRAMMIIRRMAHRFQT
	l	l		l		EEIKRISFIKRRVGNKDEINPGA
				1		RWRRFHDDGITFLRLLDAGLTI
		l				PSGKTIVHPLARQRRGLVKIQFL
			l		İ	R*LLAKLTRQPLLPLQPVTARL
			İ			CLNPRLRRQRKIRMPAGTKWQ
					l	RLCPLIGDR*RHRQRSAVYALS
		1				VRHNPVHHILRKNGDMYSFPIT
			i		1	IGNNVWIESHVVINPGVTIGDNS
		1				VIGAGSIVTKDIPPNVVAAGVP
						CRVILEINDRDK
22616	52984	Α	22744	742	1158	
22617	52985	A	22745	439	2882	PDWQAFQSPPFPASLIIHPKPAK
						LPGWLCIVQWSLLAITRTPTPV
						RWAOOTTETVGLVVGDVSDPF
		ŀ			ŀ	FGAMVKAVEQVAYHTGNFLLI
		1			ł	GNGYHNEQKERQAIEQLIRHRC
		1			ŀ	AALVVHAKMIPDADLASLMKO
		1				MPADKOCESIPDONNOTGTOY
		ı		ł	ł	PRGKEPKLLRRLMHMNRKVGP
		1				RGPRRMLLRYLDVEFLSVALA
		1				VVHLGPQGTCSRLFLVARALV
						VAVSEMDKSLSAQAQIYFASSN
		1				GPVPDLAYOVDFPRLEIGL*F\E
		l				YVDTGAETTLVPGDVLYVPAG
		1				GWNFPQWQAPATTFSVLFGKQ
		ı				QLGF/SAVVQWDGKQYQNLAK
		ŀ				OHVARRGPRIGSFLLOTLNEMO
		ŀ				MQPQEQQTARLIVASLLNHCR
						DLLGSQIQTASRSQALFEAIRDY
						IDERYASALTRESVAQAFYISPN
		ŀ				YLSHLFQKTGAIGFNENLSHTR LGHA*TLLKGYDLKVKEVAHA
		1	1	l	l	CGFVDSNYFCRLFRKNTERSPS
		1		l		EYRRHRRLHDNRICQFKTPTDN
	l	l				PENPDRYLTFRRIDKETSSLPKK
	1	l				VAPSCKRLSVSRSSPKPSISIRRC
		1				MRLMLIEGRRSSFIHPDKDVDG
		1				FHPYNVGRLCQRAPRLRPCTPR
		l		l		GIVTLLERYNIDTFGLNAVVIGA
		I		l		SNIVGRPMSMELLLAGCTTTVT
		l				HRFTKNLRHHVENADLLIVAV
		1	1	l		GKPGFIPGDWIKEGAIVMMSGI
		<u></u>				NRGVATDASCFPLVPFANRVSG
22618	52986	Α	22746	2	126	
22619	52987	A	22747	1	465	
22620	52988	С	22748	78	308	

SEQ ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletinn, \-possible nucleotide insertion)
-				sequence		
	<u> </u>	<u> </u>				
22621	52989	Α	22749	I	2797	MEHPPVLAVQRVALPSRSSEPE
	i	ŀ				LFLNPPTDEETEAHRGMYLDQ
1	1					GPRTKDWIPVAPGVREPSPLGP
1						GPSSLWLLVVGGTVEEGSRKH
						RPTCKKDFTKSKTVDQAPTLCA
						RHQMLNIQFLISDRDPQCNLHC
					ŀ	SRTQPKPICASDGRSYESMCEY
		1				QRAKCRDPTLGVVHRGRCKDA
i						GQSKCRLERAQALEQAKKPQE
		1				AVFVPECGEDGSFTQNTSCIHSS
				İ		VYQIGENSSLTFLEKREESGHL
				!		QRDWFPLWFPRKARVI
22622	52990	Α	22750	1	5295	MNLNQEEIPDLPEKEFRRLVIKI
		1		l		IREGLEKGKAQCKEIQKMIQEV
1		1		i		KREIFQEIESLKKKPPKIQKTLDI
		l		l		LLKMONALESLSDRTEOAEEIN
		l				SELEDKVFKLTQSNKDKEKITR
1		i i				KYEOSLOEVWSYVKRPNLRIIG
		l	i			VPEEEENTKSLENILGGIIEENFP
l		l		ł	1	SFARDLDIQIPEAQRTTGKLIAK
		l				RSLLRHIVFRLSKVKMKERILR
		l		İ		AVROKHOATVSOKNKAGGITL
		1				PDFKLYYKATVSQKNKAGGITL
1		1				PDFKLYYK
22623	52991	Α	2275I	1	891	GSPFEVMPRDDVFFIVDVKI*DI
1		1				GVYSCTAONSAGSISANATLTV
		1				LETPSFLRPLLDRTVTKGETAV
		ı	1			LQCIAGGSPPPKLNWTKDDSPL
		1				VVTERHFFAAGNQLLIIGDSDV
1			ŀ	İ		SDAGKYTCEMSNTLGTERGNV
1						RLSVIPTPTCDSPQMTVL/HLLD
						DDGWGHCGLYVIIAVVCCVVG
	l	l	İ			TSLVWVVIIYHTRRRNEDCSITN
						TDETNLPADIPSYLSSOGTLADR
		l				ODGYVSSESGSHHOFVTSSGAG
						FFLPQHDSSGTCHIDNSSEADVE
						AATDLFLCPFLGSTGPYVFEGK
						CVWAONPF
22624	52992	A	22752	1	2947	MSAPSLRARAAGLGLLLCAVL
						GRAGRSDSGGRGELGOPSGVA
		1				AERPCPTTCRCLGDLLDCSRKR
		1		l		LARLPEPLPSWVARLDLSHNRL
		1				SFIKASSMSHLQSLREVAGNRIV
		1	1			EILPEHLKEFOSLETLDLSSNNIS
		1				ELOTAFPALOLKYLELNRNKIK
		l				NVDGLTFQGLGALKSLKMQRN
1				l		GVTKLMDGAFWGLSNMEILQL
1		1				DHNNLTEITKGWLYGLLMLQE
1		1	1			LHLSQNAINRISPDA WEFCQKL
		1				SELDLTFNHLSRLDD
22625	52993	A	22753	2	417	
22023	02//3	l.,	22177	<u>-</u>	1	L

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
22626	52994	A	22754	I	262	VLDIGS/SLTVPDEFSKQE/RAD GMWWKQLVAGAVAGAVS/RT GTAPLDRLKVFMQIKRAILGQQ ETLMCRSASWLAPWLVPQPNH HLPYG
22627	52995	Α	22755	1	762	
222628	52996	Α	22756		1527	ILLAAGADPAGSKGGGSREA PTEAARCRASPPRAGAMRGSP GDAERRQRWGRLFEELDSNKD GRVDVHELRQGLARLGGGNPD PGAQQGISSEGDADPNGGLDLE EFSRYLOEREGRLLLMFISLDR NQDGHIDVSEIQQSFRALGISIL LEQAEKNFAQA/VDRDGTMTID WQEWRDHFLLHSLENVEDVLY FWKHSTVLDIGECLTVPDEFSK QEKLTGMWWKQLAAGAVAG AVSRTGTVPLDRILKVFMQVHA SKTYNELNILGGI-SMYLEGGIRS LWRGNGINVLKIAPESAIKFMA GSLAGATAQTINYPMEVLKTRL TLRRITGQYKGLAGLRGGILER EGPRAFYRGYLPNVLGIIPYAGI DLAVYETILKNWWLQQYSHDS APGILVLACGTISSTCQJIAS YPLALVRTRMQAQSIEGGPKL SMLGLRHILSQEGMRGLYRGT SMYLERFYAGYLPNVLGIIPYAGI SMLGLRHILSQEGMRGLYRGT SMLGRHGWGYNGYMGYMGYMGYMGYNGYMGYNGYMGYNGYMGYNGYMGYNGYMGYNGYMGYNGYMGYMGYMGYMGYMGYMGYMGYMGYMGYMGYMGYMGYMG

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		-
22629	52997	A	22757	1	2127	MDQSGLLALRAAVGAVVGSTL
						AVDIGILVKDWHPRPLGGKHC
						QLLGAFRCRQHRRGKEERRGE
						TKEKKGERENERGKRRRIEGER
						DKYVKENEKQTCVQKERHSCT
			İ			GNSPNHMATVQAERGALDSHG
						GWHPWDYCAVQQGRYHQLLL
						TKEEIEVQRSKKMCLLQAFLIC
						CQKSHFLGPGCTRSMVPASASG
						ESLKLLPLMVEDEGEPVTERE\R
1					1	KREGERERGRERERRRRKRK
						RRRKRRRKRRRRRRRG*EFLS
						LG*NERGKRRRIEGERDKYVKE
		1				NEKQTCVQKERHSCTGNSPNH
		1				MATVQAERGALDSHGGWHPW
		1			1	DYCAVQQGRYHQLLLTKEEIE
1						VQRSKKMCLLQAFLICCQKSHF
		ļ.				LGPGCTRSMVPASASGESLKLL
		į .			1	PLMVEDEGEPVTEREQEEGGGE
				ł		REREREKKEKEKEKEKEEK
						EKEKEKEEERIRVPLPRVELRLT
		ı			ì	NIAAKQVGNTILDVAAPADHPS
1						ESSCVYVRFVIKIPVAASFRFKK
		[				SYVTNNVESCEDGIHYFWKSH
		1				MHSENSCSCKLPCPPHLPAAER
		ı				LGDALEPLLPTMRRQRQCQNT
		l	ļ	i		GDCDGRRQIGTRETEMHRDTSL
ł		l				VAVKTQVDTSREGMGEVEPNS
		1				CFWRDTHMASGVGAAFEELPH
1		1				DGTCDECEPDEAPGAEEVCREC
i		1				GFCYCRRHAEAHRQKFLSHHL
1		1				AEYVHGSQAWTPPADGEGAGK
22630	52998	A	22758	348	881	WROPRWPHHOACLWORAGSS
		1			ĺ	YSOHKDPHFTSFSTPRAVONFD
		l				GFPCPFLRETTTLVLIPSVPLTDP
l		ł				GSFYRAWSFQCGPRSPRPLLGR
İ		1				TLSQKFLLRLRGRPRRPYMSTL
ľ		1				TPF*STRIFERAFCQVMGAESGG
i						RIEKTTLCKRSLGPLMEPRV
22631	52999	A	22759	I	347	MVPASASGESLKLLPLMVEDER
				1	Ī · ·	EPVTERE/REKGRGREREGERE
				1	l	KEREGEREREREGERKGERER
				1		RRG*QPPFHNRFSEGQGRSFLCS
						RRPRVSRRGHTSRGPARVKAAE
						RRGLG
22632	53000	A	22760	145	258	PENGLIGLS*KIVLSLNIPVNEN
12032	155000	l.,	/-	1	1	DMKVDEVLYEDSSTA
L			<u> </u>	L	l	DMICTODIOLOGIA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22633	53001	Α	22761	1	1479	MALKEIFCPVRETETCTINCGTR
						ISPSILHVGLPYPSTPNVKPERE
		1				GCGGGVSGNSKQPGALETAESI
		1				GKKEGRREEGSRPESASSLSLLL
		1				SLAYDVDKSAQICTRIHSSVPLE
		1			1	AVYPGYIHQSIINQADLSALAPY
						DPCVCQIHKYHGAYNCLQYSV
		1				QNTLYSPTTEQYKRCQAPWVL
	Ì	1				PAIFTIPASGAAASAVPVRAQA
		1				QGGQAAGQRRRLATAPLTGEG
	1					EPDHGDRTRAGSQGPEARPRIH
			ļ			LGAARGGDRTRGRPGRRGRDW
		1				TRLGREMAAAAVSSAKRSLRG
						ELKQRLRAMSAEERLRQSRVLS
		1		ļ		QK\VIAHSEYQKSKRISIFLSMQ
		1				DEIETEEIIKD\ICQRGK\ICFIPPV
		1				RFSRAITMDMVRIGSP\EGNFFN
		ı				FPKTSWEYSLRLGEG\DVREEG
	1					LVPPGGL/EILILPCPQALGLDQ
		I				NHGQPRLG\RGQGAYL*CLS*K
1	į.	1				RLFCSHSGKLKPPTPPGGWLFK
						EQICPPRSPVNENDMK\VDEVL
22634	53002	A	22762	1	813	
22635	53003	Α	22763	1	368	
22636	53004	Α	22764	327	4281	SASASRCDPGSGSRREREEELQ
İ		1	ŀ			WRRRRRRRRRRRRRQRQRA
1		1				AAPAAPAGGIEAVNMASASYHI
Į.		1				SNLLEKMTSSDKDFRFMATND
						LMTELQKDSIKLDDDSERKVV
						KMILKLLEDKNGEVQNLAVKC
			l			LGPLVSKVKEYQVETIVDTLCT
						NMLSDKEQLRDISSIGLKTVIGE
		1	1			LPPASSGSALAANVCKKITGRL
			l			TSAIAKQEDVSVQLEALDIMAD
						MLSRQGGLLVNFHPSILTCLLP
	ļ	Ļ				QLTSPRLAVRKRTIIA
22637	53005	A	22765	2	271	
22638	53006	A	22766	299	451	IVALFRSLVRHLRGTMAMESTA
				1	1	TAAVAAEL/VSADKIEDAPAPST
1	1	1	I			SADKVES

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22639	53007	A	22767	1	1122	MATESTATAAIAAELVSANKIE
22039	33007	<u> </u> ^	22707	ľ	1122	DVPAPSTSADKVERIENGVLGN
			İ			TLDGVHVEEEEGEKTEDESLVE
						NNDSIDEEARKELREOVYDSM
			l	i		GEKEEAKKTEDKSFAKPETDKE
						ODCEMEKGGREDMDISESAEEP
	ŀ		İ			OEKVDLTLGWLTEISEGAKGG
				i		GAPEGPNEAEVTSGKPEOEVPD
						VEEEKSVSETDVQKECREKGG
						WEKHREVIVSIEEKPKEVSEEOP
						VVILEKQGTAVEVAAESLDPTV
l						KPVDVGGDEPEEK/GTVPGSLR
		1		1		
1		1			1	PSPCRDPLPAGLGLWVQLSV**
				ł		GSGTVQQIY*SH*EENGCTKRA
		1				GEGG*RIVC*IQERN*GTEGTAT
		1		1		RNYRKDRRCKGVSDYWECS*T
22640	50000	١.	007/0	222	339	GSESYSGGELYFRFHS*WRRLF
22640	53008	A	22768	223	496	
22641	53009	В	22769	12	1099	VSGVPNCLFFSOALFRSLVRHL
22642	53010	A	22//0	12	1099	
		1				RGTMAMESTATAAVAAELVSA
			1			DKIEDVPAP\STS\ADKVESLDV
		1			ŀ	DSEAKKLLGLGQK\HLVMGDIP
				l	İ	AAVNAFQEAASLLGKKYGETA
		1				NECGEAFFFYG\KSLLELARME
		l			l	NGVLGNA\LEGVHVEEEEGEKT
1		l				EDESLVENNDNI/D*N*RLGRG*
1						*RK**D*RNAK*FSP*KQVSSRK
		1				*GGGDWEPRACLGYAGFSKDH
Ì		1				F*KARNKRSTALCCPGTS*TRRS
					1	*C*I*\KLCASC\GGVPVLP*PTGP
		1	1	i	1	VPGSPRPSPCRDPLPAGLGLWV
		l				QLSV**GSGTVQQIY*SH*EQNG
		l			l	CTKRAGEGG*RIVC*IQERN*GT
		İ				KGTATRN*REDRRCKGVSA*W
1		l		L		ECS*TGSESYS

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
		<u> </u>			Ļ	
22643	53011	A	22771	11	2485	VSGVPNCLFFSQALFRSLVRHL
						RGTMAMESTATAAVAADVVS
						ADKIEDVPAPSTSADKVESLDV
						DSEAKKLLGLGQKHLVMGGIP
			l			AAV\NAFQGAA\SLL\GKKYGET
						ANE\CGEAFFFYG\KSLLELARM
	i		İ			ENGVLGNALEGVHVEEEEG\EK
	1					TEDESLVENNDN\IDEEAREELR
						EQVYDAMGEKEEAKKTEDKSL
	1					AKPETDKEODSEMEKGGREDM
1						DISKSAEEPQEKVDLTLDWLTE
						TSEEAKGGAAPEGPNEAEVTSG
	1					KPGQ\EVPDAEEEKSVSGTDVQ
						EECREKGGOEKOGEVIVSIEEK
						PKEVSEEOPVVTLEKOGTAVEV
					ľ	EAESLDPTVKPVDVGGDEPEEK
						VVTSENEAGKAVLEOLVGOEV
	1					PPAEESPEVQTEAAEASAVEAG
						SEVSEKPGQEAPVLPKDGAVN
		l				GPSVVGDQTPIEPQTSIERLTET
	i					KDGSGLEEKVRAKLVPSQEETK
	İ		l			LSVEESEAAGDGVDTKVAQGA
						TEKSPEDKVQIAANEETQEREE
1						QMKEGEETEGSEEDDKENDKT
1						EEMPNDSVLENKSLQENEEEEI
						GNLELA WDMLDLAKIIFKRQET
		l	i .			KEAQLYAAQAHLKLGEVSVES
	ĺ					ENYVQAVEEFQSCLNLQEQYL
			İ			EAHDRLLAETHYQLGLAYGYN
						SQYDEAVAQFSKSIEVIENRMA
				i		VLNEQVKEAEGSSAEYKKEIEE
				1		LKELLP\EIREKIENAKE\SQR\SG
22644	53012	Λ	22772	52	161	FONPTGLY/CDTEYN\RLOKOO
			_			TEKTKOKKTOFHPICN
22645	53013	A	22773	3	115	FFQNPTGLY/CDTEYN/RLQKQQ
		1		1		TEKTKOKKTOFHPICN
22646	53014	A	22774	229	358	
22647	53015	A	22775	257	386	
22648	53016	A	22776	259	294	
22649	53017	A	22777	217	398	KAWWNLLRKSEPKSCGS*GGL
22049	33017	l^	22///	1217	370	STSRERLVECSSPLGLGYCG**L
1	1	1	l	l		CHLOEPHYGSLHRMSS
22650	52010		22770	l	2204	CHEQUETT I GSETIKMSS
22650	53018	A	22778	11	3386	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22651	53019	A	22779	391	2100	SRRTAGVRLDILCKDPIMWVSA
						DGHSYQPQYPKAKRATEFHFF
						NFKALLARAAVGAARGIHIHRC
		l				GLTKRPDARRRRVAGAOGREP
				i		AISLPGDRAAGARATRTRGPGP
		ĺ				APKMPAIAVLAAAAAA\WCFL
1						QVESRHLDALAGGAGPNHG\NF
						L\DNDQ\WLSTVSQYDRDKY\W
						NRFRDEVEDDYFRNW\NPNKPF
						DQAL\DPSKDPCLKVKCSPHKV\
						CVTQ\DYQTAL\CV\$RKHLLPR
						QKEWGTWAQKHWGLDL*ILV
						K\CKPCSRGRSSAMGLAGSDGP
	l					LLTAFQVANLEFHACSTGQK/S
	ŀ		į.			FATLCDG\PCPCSSQSLEPPKHK
					ļ	GRKGVPCTDKELRNLASRLKD
						WFGALHEDANRVIKPTSSNTAQ
	İ				1	GRFDTSILPICKDSLGWMFNKL
					i	DMNYDLLLDPSEINAIYLDKYE
			•			PCIKPLFNSCDSFKDGKPFLNNE
						WCLLPSQNPGGLP/CAQNEMNR
		1				IQ\KLSKGKSLLGAFIPRCNEEG
		l			ľ	YYKATQCHGSTGQCWCVDKY
1						GNELAGSRKQGAVSCEEEQETS
1						GDFGSGGSVVLLDDLEYERELG
						PKDKEGKLRVHTRAVTEDDED
						EDDDKEDEVGYIW
22652	53020	A	22780	2	441	HGVQGAGCGEVAAVSIELGAG
						GPVWRAAPRACSRLQLLRGFG
						LEGQWFGESCCRRGAS*EGSTR
						*PPPRRQPWPEWVMSWVPPET
				ŀ		REPARISPAPAKQN*PPPAPKSP
1						AAPA\PQAAVAATPSPPGEISEA
		_				WDSLESFSKPAFFSQ
22653	53021	Α	22781	3	121	

SEQ ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hed	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22654	53022	Α	22782	2	1488	FVNNMADGEEPEKKRRRIEELL
					ľ	AEKMAVDGGCGDTETWKDR\
						WNHVKK\FLERSGPFTHPDFEP\
						STESLQFLLDTCK\VLVIGAGG\
						LGCELLKNLALS\GFRQIHVID\
						MDTI\DV\$N\$*IRQFLFRPKDIGR
						PKAEVAAEFLNDRV\PNC\NVV
		l				PHF\NK\IQDFN\DTFY\RQFHI\IV
						CGTGTLSCARRWINGMLISL\LN
			l			YERMVS*DPSSIVPLID\GGTEGF
						\KENARVIL\PGMTACIECTLEL
						YPPQVNFPMCTIASMPRLPEHCI
						EYVGMLQWPKEQPFGEG\VPL
						DGDDP\EHIQWIFQKSLERASQY
						NIRGVTYRLTQGVVKRIIPAVA
						STNAVIAAVCAT*/EVFKIATSA
			i			YIPL*IITWVFNDV\DGLYTYTFE
						GRKGK\ENCPAC\SQLPQNISVF
						LHQAKLQ\EVLGIILTNSASLQN
		i				EILPAITATLGGEKIGTLYLQSV
						TSI\EERNRANFFQTIERNWGLL
1		1				DGPKTGRLLDVTTP\QTVLFQTS
				1		ILLLKGKSPT
22655	53023	A	22783	1	1719	
		_				
22656	53024	В	22784	199	460	
22656	53024 53025	B A	22784 22785	199 226	460 1620	NGSGLALATGNRDPGLCPLSAP
		_				SCPWR*KSAGASTQQAYSSQSQ
		_				SCPWR*KSAGASTQQAYSSQSQ /HPAYHRITPT\LTNFALRLYKEL
		_				SCPWR*KSAGASTQQAYSSQSQ /HPAYHRITPT\LTNFALRLYKEL AADAPGNIFFSPVSIS\TTLCLLC
		_				SCPWR*KSAGASTQQAYSSQSQ /HPAYHRITPT\LTNFALRLYKEL AADAPGNIFFSPVSIS\TTLCLLC LG\AQANNSALILEGLGFNLTET
		_				SCPWR*KSAGASTQQAYSSQSQ /HPAYHRITPTLTNFALRLYKEL AADAPGNIFFSPVSIS\TTLCLLC LG\AQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL AADAPGNIFFSPVSISTTLCLLC LGVAQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL ELKVGNSLFLDKRLKPRQHYFD
		_				SCPWR*KSAGASTQQAYSSQSQ #PAYHRITPTLTNFALRLYKEL AADAPGINFSPVSISTTLCLLC LGVAQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSFKL ELKVGNSLFILDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGR
		_				SCPWR*KSAGASTQQAYSSQSQ /HPAYHRITPTLTNFALRLYKEL AADAPGNIFFSPVSIS\TTLCLLC LGYAQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL ELKVGNSLFLDKRLKPRQHYFD SIKELYRAFASAHFTDSVTTGR QINDYL*RRTYWPGVDLPFRS/L
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLINFALRLYKEL AADAPONIFFSPVSISITILCLLC LGWQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL ELKVGNSLFLDKBLKFRQHYFD SIKEL YRAFAFSAHFTDSVTTGR QINDYL*RRTYWFGVDLPFRS/L ARDTFMVLANFILESKAKVKA
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL AADAPONIFSPVSISITTLCLLC LG\(\)QANNSALILEGLGFNLTET PEADIHQGFRSILHTLVLPSPKL ELK\(\)GNSILHTLVLPSPKL ELK\(\)GNSILDKRLFRQHYFD SIKELYRAFAFSAHFTDSVTTGR QINDVL*RRTYWFGVOLPFRS/L ARDTFMVLAINFILFSKAK\(\)KA TLSSRVPDPKSRPVFLVDERTSF
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL AADAPONIFFSPVSISITTLCLLC LGXQANNSALILEGLGFNLTET PEADIHQCFSSLLHTI-VLSPKL ELKVGNSLFI.DKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGR QINDVL*RRTYWFGVDLPFRS/L ARDTFMVLAINFILESKAKVKA TLSSRVPDPKSRPVFLVDERTSF QVPMMHQKEMHRFLYQQDLA
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL AADAPGNIFFSPVSISITTLCLLC LGYQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL ELKVGNSLFLDKRLKFRQHYFD SIKEL YRAFAFSAHFTDSVTTGR QINDYL*RRTYWFGVDLPFRS/L ARDTFMVLANFILESKAKVKA TLSSRVPDPKSRPVFLVDERTSR QVPMMHQKEMHRFLYDQLLA CTVLQIESYRGNALALLVLPDFG CTVLQIESYRGNALALLVLPDFG
		_				SCPWR*KSAGASTQQAYSSQSQ #PAYHRITPTLTNFALRLYKEL AADAPONIFFSPVSISTTLCLLC LGXQANNSALILEGLGFNLTET FEADIHQGFSSLLHTLVL.PSPKL ELKVGNSLFLDKRLKPRQHYFD SIKEL YRAFAFSAHFTDSVTTGR QINDYL*RRTYWFGVDLPFRS/L ARDITFMVLANFILFSKAKVKA TLSSRVPDRSKRPVELVDET LOYDMMHQKEMIRFLYDQDLA CTVLQIELYRGNALALLVLPDFG KMKQVEAALQPQTLRKWGQL
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL LGAQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL ELKVGNSLFLDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGR GINDVL*RRTYWGVOLDFRS/L ARDTFMVLAINFILFSKAKVKA TLSSRVDPKSRPVLDERTSF QVPMMHQKEMHRFLYQOLLA CTVLQIEVRGHALLVLPPG KMKQVEAALQPQTLRKWGQL LLPSLLDLHPRFSISGTYNLEDI
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL AADAPONIFFSPVSISITTLCLLC LGQAQANNSALILEGLGFNLTET: ELKVGNSLFILDKRLKPRCHYEP ELKVGNSLFILDKRLKPRCHYEP ELKVGNSLFILDKRLKPRCHYEP GVINDYL*RRTYWPGVDLPFRS/L ARDITMVLANIFILESKAKVKA TLSSRVPPRSRPVFLVDETST QVPMMHQKEMHRFLYDQDLA CTVLQIEVTRGMALALLYLDPE KMKQVEAALQPCITLRKWGQL LLPSLLDLHLPRSISGTYNLEDI LEPSNWFSPTYTLNKITSLQSVG
		_				SCPWR*KSAGASTQQAYSSQSQ #IPAYHRITPTLTNFALRLYKEL AADAPONIFFSPVSISITTLCLLC LGQQANNSALILEGLGFNLTET FEADIHQCFSSLLHTI.VL.SPSKL ELKVGNSLFLDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGR GINDYL*RRTYWFGVDLPFRS/L ARDTFMVLANFILESKAKVKA TLSSRVPDPSSRPVELVDETT QVPMMHQKEMHRFLYDQDLA CTVLQIEVRGNALALLVLPDFG KMKQVEAALQPOTLRKWGQL LLPSLLDLHLPRFSISGTYNLED LSPNWFSPTYTLNKITSLQSVG G*LNKSISKVSHKSMVDMMEK
		_				SCPWR*KSAGASTQQAYSSQSQ APPAYHRITPTLINFALRLYKEL LGAQANNSALILEGLGFNLTET PEADIHQGFRSILHTIVLPSPKL ELKVGNSI-FLDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGR GINDVL*RRTYWGFOULPFRS/L ARDTFMVLA\NFILFSKAKVKA TLSSRYDPKSRPVFLVDERTSF OVPMMHQKEMHRFLYDODLA CTV-LQIEVYRGNALALLVLPDPG KMKQVEAALQPQTLTR.WGQL LLPSLLDLHPRSISGTYNLEDI LSPNWFSFTYTLNKITSLQSVG G*LNKSISKVSHKSMVDMNEK GDQRPGLLQASSPSPHL*YTIMS
		_				SCPWR*KSAGASTQQAYSSQSQ #IPAYHRITPTLTNFALRLYKEL AADAPONIFFSPVSISTTLCLLC LGXQANNSALILEGLGFNLTET EEADHQGFRSELHTI-UL-PSPKL ELK-VGNSLF-LDKRLK-PR-QHYPE SIKEL YRAFAFSAHFTDSVTTGR QINDYL-PKRTYWFGVDLPFRS/L ARDITFMVLANIFILFSKAK-VKA TLSSR-VPDFSSRPVFLVDSETS- (VDFWMHQKEMIRFLYDQDLA CTVLQIELYRGNALALL-UL-PDFG KMKQVEAALQPQTLRK-WGQL LLPSLLDLH-PFFSISGTYNLEDI LSPNWFSFT/TLNKIFISLOGG G*LNKSISK-VSHKSMYDMNEK GDQRPGLLQASSPPL*-YTMS DPHAHFINRPFLLLL-WEVTQSL
		_				SCPWR*KSAGASTQQAYSSQSQ APPAYHRITPTLINFALRLYKEL LGAQANNSALILEGLGFNLTET PEADIHQGFRSILHTIVLPSPKL ELKVGNSI-FLDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGR GINDVL*RRTYWGFOULPFRS/L ARDTFMVLA\NFILFSKAKVKA TLSSRYDPKSRPVFLVDERTSF OVPMMHQKEMHRFLYDODLA CTV-LQIEVYRGNALALLVLPDPG KMKQVEAALQPQTLTR.WGQL LLPSLLDLHPRSISGTYNLEDI LSPNWFSFTYTLNKITSLQSVG G*LNKSISKVSHKSMVDMNEK GDQRPGLLQASSPSPHL*YTIMS
		_	22785	226	1620	SCPWR*KSAGASTQQAYSSQSQ #IPAYHRITPTLTNFALRLYKEL AADAPONIFFSPVSISTTLCLLC LGXQANNSALILEGLGFNLTET EEADHQGFRSELHTI-UL-PSPKL ELK-VGNSLF-LDKRLK-PR-QHYPE SIKEL YRAFAFSAHFTDSVTTGR QINDYL-PKRTYWFGVDL-PFSXLAKVKA TLSSR-VPDPKSRPVFL-VDST-VGVPMMHQKEMHRFL-YD-QDL-A CTVLQIE-YRGNALALL-VL-PDFG KMKQVEAALQPOTL-RK-WGQL LLPSLLDLH-PFFSISGTYNLEDI LSPNWFSPT/TLNKITSLGVG G*LNKSISK-VSHKSMYDMNEK GDQRPGLLQASSPPL*-YTMS DPHAHFINPFLLLL-WEVTTQSL
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL LGAQANNSALILEGLGFNLTET ERADHQCPSSELLHTI.VLSPKL ELKVGNSLFLDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGK GINDTL*RRTYWFGVDLPFRS/L ARDTFMVLAINFILESKAKVKA TLSSRVPDPKSRPVFLVDERTSF QVPMMHQKEMHRFLYDQDLA CTVLQIEVRGNALALLVLPDPG KMKQVPAALQPQTLRKWGQL LLPSLLDLHLPRFSISGTYNLEDI LSPNWFSFTYTLNKITSLQSVG G*LNSISKYSKHSMVDMMEK GDQRPGLLQASSPSPHL*YYTMS DPHAHFNRPFLLLLWEVTTQSL LFLGKVVNPVVWTMVGQQE

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22660	53028	A	22788	309	2286	CAVCHISFQDTPVLVSSNVTMQ
		l				FGSKPLFENISVKFGGGNRYGL\
						IGANGSGKSTFMKILGGDLEPT
		1				LGNGSLDPNER\IGKLRQ\DQFA
		1				FEEFTVLDTVIMGHKEL\WKVK
	ŀ					QERDRIYALPEMSEEDGYKVA
	1	1				DLEVKYGEMAGYSAEARAGEL
					1	LLG\VGIPVEQHYGPMSEVAPG
	ł	İ			l	WKLRVLLAQALFADPDILLLDE
						PTNNLDIDTIRWLEQVLNENNS
		1				T\MIIISHDRHFLNMVCTHMAD
						LDYGELRVYPGNYDEYMTAAT
					1	QARERLLADNAKKKAQIAELQ
		l				SFIDKIKLEEVKASSRQNPFIRFE
						QDKKLFRNALEVEGLTKGFDN
	1				ł	GPLFKNLNLLLEVGEKLAVLG'T
						NGVGKSTL\LKTLVGNLQPDSG
	1				l .	TVKWSENARIGYYAQDHEYEF
						ENDLPVFEWMSQCKQ*GDDEQ
						AVRSIFGRLLFSQDVIKKPAKVL
						SAPFCKDSILEAIDAGIKLIITITE
	1				1	GIPTLDMLTVKVKLDEAGVRMI
	1			l	l	GPNCPGVITPGECKIGIQPGHIH
	1					KPGTGELNITTGGIVKARDTQIA
	1			ļ		LNDKSKGDVRVDGQNSLLETF
						NMYVGTSGTGTLTLTNNGTLN
	1	l				VEGGEVYLGVFEPAVGTLNIGA
	i	ŀ				AHGEAAADAGFITNATKVEFG
	1	l				LGEGVFVFNHTNNSDAGYQVD
						MLITGDDKDGKVIHDARHTVF
22661	53029	Α	22789	I	553	
22662	53030	Α	22790	1633	2207	ILFFDKGMPAQTPSLQKRWEK
	i	1				M*LVLW*GDREGIKAI/ISEESW
		l	l			GARAPIFLLR/MMEDQGLYSIR
		ı				MGGARGLPGKSLAL*GAGRGF
						YSGERTWTLSLPSGNCAP\YQR
						GGWWYHACA\HSNLNGVWHH
						GGHYRSRYQDGVYWAEFRGG
				l		AYSLRKAAMPHLAPEAVTLCSF
		_				GPLGPRGHWSAGAPSCSGHTFF
22663	53031	Α	22791	34	210	AAADPLRHLRRGAFIKAGRHA
			l			G/PSLPVCLQATA\H*MDLARLA
	<u> </u>					LVLVVRTSASTKSSTRE

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22664	53032	l la	22792	397	1168	LLSRISKVSSKLPVRGGAGSLCT
						GDQRK\RRKFL\ETVELQISLKN
						YDPOEGOSAFSGHPSRL*V\PSP
		l				PLSSLVCVLGGPSKHL*RRLRP
		1				VGYPPTMDIEGA*KKLNOELKN
		İ				LVKKAGPRKYDCVFWPQESLD
		ı				SSKIPPNPSGPGFK*RAGKVPFP
		l				CLTHNGKHGLAQSGIEVEVPTF.
		l				LSFQMKKGVYVWAVAVWSRE
		l				R*QTNELVYNIHLACQLSWVSL
		1				LKEKNWQ\NVPGPYIIKSTMGQ
						SPSSHILRHIRINSITS
22665	53033	Α	22793	1	377	SVSRVPTRSLRTSTWRLAQDQT
		İ				QDTQLITVDEKLDITTLTGVPEE
		l				HIKTRKVRIFVPARNNMQSGVN
		l				NTKKWKMEFDTRERWENPLM
		l				GWAST\WSYDIEERKVPKPKSK
22555	50004	Ļ.	00004	100	260	SYGANFSWNKRTRVSTK
22666 22667	53034	A	22794	183	368 623	ILAFACSKMAAVSMSV\VLROT
22007	33033	^	22193	ľ	023	LWRRRAVAVAAL\SVSRVPT/R
		l		1		RSLRTSTWR\LAH\DQTQDTQLI
		l		i		TS**KIGISLTFNWPFPEEHIKNL
		l	i	l		EKSGIFVPCSPITCOSGVNNTK\
		l		l		KWK\MEFDT\RER\WEN\PLMG\
				ŀ		WASTGLIPLS\NMVLNLPVLKE
		ł		l		DAVS\FAEK\NGWELWTLEERK\
		ł				VPKPKSQVFMGANFS\WNKRTR
		1		1		VSTKIGLALTISPA
22668	53036	Α	22796	1	798	MNKVNPTQQDKEDHPELIITSIG
		l		1		DKRQLTLEAASGVCHRGTQTR
		1				VKEESKKFLISMSFRGGKPVHK
		l		1	İ	VVERACVCTCLHHTSVPATHN
		l		l		TQRRHTTPTTRKNGRKRRGKK
		l		l		EREKRKDREHNTGRRTKRKKT
		l				RNIGGQRTQQKQKQSEDKARR
				l		EPGGGRKQGRGREDGTRQRRG
		1	l			GRTRNREEGAQRGRNGR/RGEG
		1	l	l		RRTEEETHMA*APDITATQQTE
			l			QQGHGTRQRDTTLRRHTAEDA
						QPSEHASADQAHTKKTE*SVSR
22669	53037	В	22797	85	322	TNGQYLLT
22009	22027	ľ	44171	102	322	

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codnn for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			}	sequence		
22670	53038	A	22798	165	4733	AATALLTHGQEEGQVEGQDED
		1				IPPITCVQNGLRYHDRDVWKPE
						PCRICVCDNGKVLCDDVICDET
		1		1		KNCPGAEVPEGECCPVCPDGSR
		1		l		VTHRPKKPPGVEGPKGDTGPPR
		1				PKGT\PGPPGRDGIPGQPGLPGP
		i		l	ŀ	PGPPGPPGPPGLGGKLLLPKLSY
		i			-	GYDEKSTGGISVPGPMGPSGPR
	l			1		GLPGPPGAPVSIQDVFICLLGLW
	1				ŀ	SFGGKTGMRAGEMLRDLLGPO
				l	ŀ	GFQGPPGEPGEPGASGPMGPRG
	1			l	İ	PPGPPGKNGDDG
22671	53039	С	22799	88	392	PFGFFGKNGDDG
22672	53040	Ā	22800	37	293	
22673	53041	A	22801	55	1917	GGGGVRRRRGTRRSEPGDSRSA
22075	15501.	ľ.		-	1	MAPIGLKAVVGEKIMHDVIKK
						VKKKGEWKVLVVDQLSMRML
						SSCCKMTDIMTEGITIVEDINKR
				i	!	REPLPSLEAVYLITPSEKSVHSLI
					i	SDFKDPPTAKYRAAHVFFTDSC
						PDALFNELVKSRAAKVIKTLTEI
	l	1				
	l	l		1		NIAFLPYESQVYSLDSADSFQSF
		l		İ		YSPHKAQMKNPILERLAEQIAT
	l					LCAT\LKEYPA\VRYRGEYKDN
	l	l				ALLAQLIQDKLDAYKADDPTM
	l			·		GEGPDKARSQLLILDRGFDPSSP
			l			VLP*I*LFQGMSY*SCLPI\ENDV\
				1		YKYETSGIGGGHGVKGGASGT
	I	1			1	EDDDLW\IALR\HKHIAEV\SQ\E
	1	1				VTRSLKDFSSSKRMNTGEKTT
	İ		ĺ			MRDLSQMLKKMPQYQKELSK
	1					YSTHLHLAEDCMKHYQGTVDK
						LCRVEQDLAMGTDAEGEKIKD
	1		l			PMRAIVPILLDANVSTYDKIRIIL
						LYIFLKNGITEENLNKLIQHAQI
	1					PPEDSEIITNMAHLGVPIVTDST
	1					LRRRSKPERKERISEQTYQLSR
						WTPIIKDIMEDTIEDKLDNK\HY
				ĺ		P\YISTR\$\$A\$F\$TTAV\\$ARYGH
	1	l		l		WP*GTRPPGEYRSGP/RVFIIFHP
						LGGVSLNEMRCALRR*PRANG
		l				KWGGA**GSTHILOPHRNLPGH
					L	E GOTT GOTTING OIT

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last eodon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, +=Stop codon, /-possible nucleotide deletion, \=possible nucleotide insertion)
22674	53042	A	22802	46	1931	GGGGVRMLCGTRRSEPGDSRS
						AMAPIGLIAVVGEKIMHDVMI.
	i					NDLNMGEWKALLVDQLSMRM
!	1					LSYCCLMTDIMTEGITIVEDINK
		l				RR*PLPSLEAVYLITPSEKSVHS
						LISDFKDPPTAKYRAAHVFFTD
						SCPDALFNELVKSRAAKVIKTL
	l					TEINIAFLPYESRVYSLDSADSF
	ŀ					QSFYSPHKAQMKNPILERLAEQ
						IATLCATLKEYPAVRYRGEYKD
						NALLAQLIQDKLDAYKADDPT
						MGEGPDKARSQLLILDRGFDPS
	l					SPVLHELTFQAMSYDLLPIEND
	ŀ					VYKYETSGIGEARVKEVLLDED
i	i					DDLWIALRHKHIAEVSQEVTRS
		1				LKDFSSSKRMNTGEKTTMRDL
						SQMLKKMPQYQKELSKYSTHL
						APCLRTCMKHYQGTVDKL\CR
						V\EQDLAMG\TDAEG\EKI\KDP
ł						MRAIVPILLDANVQTY*QIRIILL
ŀ						YIFLKNGITEENLNKLIQHAQIP
						PEDSEIITNMAHLGVPIVTDSTL
				1		RRRSKPERKERISEQTYQLSRW
		i				TPIIKDIMEDTIEDKLDTKHYPYI
	1	1			1	STRSSASFSTTAVSARYGHWHK
		1				NKAPGEYRSGPRLIIFILGGVSL
		l				NEMRCAYEVTQANGKWEVLIG
l	1	l			1	STHILTPTKFLMDLRHPDFRESS
						RVSFEDQAPTME
22675	53043	Α	22803	2	51	
22676	53044	A	22804	236	451	LHLPLFLGAIRSGQYGLWNLFR
		1				VLGWTCSLGIYLGWRVS*NFG
						RGTKALKVLSTIS/E*RNHHCHP *NNIFVY

SEQ ID NO:	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence	liou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22677	53045	A	22805	3	1173	GAGAPAAGQAGLTSTSAVRAR
		l				PVSPARQSQDPAASLSLHLSLPP
l	1					APGAGCARWRRRGPQAAMAA
						GAGAGSAPRWLRALSEPLSAA
						QLRRLEEHRYSAAGVSLLEPPL
						QLYWTWLLQWIPLWMAPNSIT
	i					LLGLAVNVVTTLVLISYCLTAT
						EEAPYWTYLLCALGLFIYQPLD
		ļ				AIDGKQARRTNSCSPLGELFDH
		1		l .		GCDSLSTVFMAVGASIAARLGT
		i i				YPDWFFSRSFYLGCLVFYC\AH
		l	l			WQTYVS\GMLRFGKVDVTEIQI
		l				AL\VIVFVL\SAFG\GATM/WGD
	1	l				YTIPILEIKLKILPVLGFLGGVIFS
		l		l		CS\NYSHVILH\GGVGKNG\STIA
		l		l		GTSVLSPGTPH/MGLIIIL\AIMIY
		l				*KVRQLMVF*KSIPCL*YPQCLG
						CVF\AKVSQKISGSSHDQK
22678	53046	A	22806	2	1091	CPGRRRRRRPGRSRARATALPT
ļ		1				PGPRKRGGPAGRKRERPERCSS
		1				SSGGSSSSGDEDGLELDGAPGR
	1	1				GKRAAQPVAAGKVGGAAVVV
		Į.				AEPEHTKKHVKLETLLITTEGK
		l			1	LWSWGRNEKAQLGHGDTKRV
	1	l				EAPRLTEGLSHEVIVC\TCARNH
		l				TLALTETGSVFAFGENKMGQL
		l			1	GLGN*TDAVPSPVQIMYNSLPIT
		l				KMA/CGYGWLGYTEQKDEMV
						PRLVKLFDFPGHGASQIYAGYT
						CSFAVSEVGGLFFWGATNTSHE
		1			1	STMYPKAVQDLCSWRIWSLAC
					i	GKSSIIVAADESTISWGPSPTFG
		l			1	ELGYRDHKPKSSTAAQEVKTL
		l			1	HGIFSEPVAMGYSHSLVIARDE
		<u> </u>				SETEKEKIKKLPEYSPQTL

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22679	53047	A	22807		1409	INSUGLASERU WAGLGLPALPI ADEGGECGETLFVWRSQKLEG SKCKÖQLLIFGATNVDLIGRKE VPKQQAAVRNLGQNLWGPHR YGCLAGVRWTTVYSGSCAAHS LLITTEGKLWSWGRNEKGQLG HGDTKRVEAPRLIEGLSHEVIVS AACGRNHTLALTETGSVFAFGE NKMGQLGILGNQTDAVFSPAQI MYNGQPITKMACGAEFSMIMG LQRKPLFLWGALNMGQLGTLQH RWAGKPIFKMACGAEFSMIMG LQRKPLFLWGALNMGQLGTCLYPR RVAIFIEKTKDGQILPVPNVVR DVACGANHTLVLDSQKVFSW GFGGYGRIGHAEQKDEMVPRL VKLFDFFGRGASQLSCWLHTC SLCQCYNGWSVFLGGHQHLPI YHVPQKQCRTSAAGESGTLAC GKSSILVAADESTISWGPSPTFG GLGYGDHRWSSTAAGEVKTL DGIFSEQVAMGYSHSLVIARDE SETEKEKIKKLEPYNPRL
22680	53048	С	22808	277	518	
22681	53049	Α	22809	2	192	
22683	53050	A	22810	258	2946	MAEAGLRGWLLWALLIRLAQ SEPYTTIHQPGYCAFYDECGKN PELSGSLMTLSNVSCLSNTPAR KITGDHLILLQKICPRLYTGPNT QACCSAKQLVSLEASLSITKAL LTRCPACSDNFVNLHCHNTCSP NQSLFINVTRVAQLGAGQLPAV AVAEAFYQHSFAEQSVDSCSRV RVPAAATLAVGTMCGVYGSAL CNAQRWLNFQGDTGNGLAPLD ITFHLLEPGQAYGSGIQPLNEGV ARCNESQGDDVATCS
22683	53051	A	22811	258	362 4038	MAEAGLRGWLLWALLLRLAO
22004	33032		2.2012	1		SEPYTTHOPGYCAFYDEGGKN PELSGSLMTLSNVSCLSNTPAR KITGDHILLICKNICPRLYTGPNT QACCSAKQLVSLEASLSITKAL LTRCPACSDNFVNLHCHNTCSP NQSLFINVTRVAQLGAQQLPAV VAYEAFYQHSFARQSVDSCSRV RVPAAATLAVGTMGGVYGSRA CNAQRWLNPQGDTGNGLAPLD ITFHILEPGQAVGSGIQPLNEGV ARCNESQGDDVATCS

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
22685	53053	A	22813	198	380	ASSESDSTPCR*MAYSWPPNSN
		l				SSHCIKELPLFMGKIPAAS*DGG
		1				PQGLAIFPPGVIVGI
22686	53054	Α	22814	1	154	
22687	53055	Α	22815	35	486	PARRGRDEC*VLAPFRATRPPS
		l				RRTR*SKATYPTPSNDQTRTIRT
		l				SRERASMPRSAEPKRPTIPRLHP
						SPRGIVTVSS/RPRSHALPVSHIH
						SCHSRTAISDNQVPAFLQVYGE
						EVRKRAGEVTQAITGFPRYRLF
						FKMLRCHRSIQVTAST
22688	53056	Α	22816	212	365	RCRSGPAPRSAPGARPASPSPPA
		1				AGTCR*RPAGHCPAGAAGLLSL
		1				CPRCST
22689	53057	Α	22817	293	618	QQNPTNCLPPASR*SKT*RARST
						AGPTRPHLHHPAAPAPGSPACA
1		l				GEGRLQR/HSPNPI.YFNWKKQR
1		1				FAAKPHFRRPQSSLNPQQWLLA
						GLLQEESSSLGTDTAPLHSLG
22690	53058	Α	22818	3	1129	FCLDDLSSAINEHLSARGRRRRS
						IVKVSLHPAVIGLKSKFLKKPD
i						QLRKLF\IGGLSFETTDESLRSHF
		1				EQ\WGTLTDCVVMRDPNTKRS
		1				RG\FGFVTYATVEEVDAAMNA
		1				RPHKVD\GRVV\EPKRAVSREDS
		1				QRPGA\HLTV\KKIFVGGIKEDT
		1				EEHHLRDYFEEIILNSMGKLEV
		1				VEIH*LDR\GQWPRKRGFAFVT
		ı				FDGHDSVDKIVIQKYHTVNG\H
1		ı				NCEVRKALSKQEMASASS\SQR
		ı			İ	G\RVGSGNFWCGRT\NGFGGND
		1			1	NFG\RGGNFSWSVVAFGG\SRG
		1				GVWIWVAVGDGYNGFG\NDGS
1		ı				HFG\GGGSY\NDFGDFH\NQSSK
1		1				FWTP*RGGNLGGQKLWPPMGG
1		ı				GG\QYFAKPTKPKVAYGRFPAA
		L				ASSYGKWPEDFN
22691	53059	Α	22819	[1	568	MGKMSPGHVRDFHGSCHHNRP
l		ı				GGPGGKSGFMGQALGPG/GCV
		1	1	1	l	QPRDLVLCVSATPAMAERGQC
			1			GAQAVASEGGSPRPWQLPCGV
						EPADFRRYMETPGCPGRNL/PA
		1		1		GAGPL*RTSAGAVWKGNVRSE
			1			SP/PQSLYWGTASGAVRRGPPSS
						RNQNGRSTDSLHCMPGKDTQ\P
		_				NP*KQPGGRLHPAKPQGRSCP
22692	53060	Α	22820	232	389	

22693   53061   A   22821   3	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
22693   33061   A   22821   3   401   VGVCEKRGAPGIPPLAPG   PVAAAVPPGPGAARAPM   AAAGGRGPPEQ*T**KAW   GTGSLJRGHPAPM**PLLR.   PLHPLPPPHEPPRPRAAISN   KCRTAAGTRQWGGRQSC   QCLR	NO:						
22693   33061   A   22821   3   401   VGVCEKRGAPGIPPLAPG		sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
PVAAAVPP/GPGASRARP   AAAGGRPEQ*T*FAXW   GTGSL/RGHPAP*PLLR.   PLHPLPPPHPLPRAAJSN   KCRTAAGTRQWGGRQSC   QCLR   QCLR   GCL					sequence		
PVAAAVPP/GPGASRARP   AAAGGRPEQ*T*FAXW   GTGSL/RGHPAP*PLLR.   PLHPLPPPHPLPRAAJSN   KCRTAAGTRQWGGRQSC   QCLR   QCLR   GCL	00.000		<del>  -</del>	Inner	10	I do 1	LIGUIGEN DO LEGISTRY LEGISLES
AAAGGRPEQ*T*FAXW	22693	53061	A	22821	3	401	
GTGSL/RGHPAPM*PLLR.			ı				
PLHPLPPPHPLERRAALSN   KCRTAAGTRQWGGRQSC QCLR							
CRTAAGTRQWGGRQSC   CQLR							
QCLR			ı				PLHPLPPPHPLPRRAA\SNPGTS
22694   53062   A   22822   1   410   MGKMSPCHVRDFHGRES   RGPGGKSGYGKVQGPH   LGTWCLVSQPLHFIG*KG   SGVASEGGSPKPWQLPR   MGAQKSRTEVKPEPLPRE   GNTWMPRIPMCCARDPG   GCSNPSQRR   GRPHADPANYSARNLSRE   GRVQDPRQAPQLPPGLL   VRRPPRRPVDRTLPGGP   AAAGPTGGRGSPRAGG   TGPRAGFQAKGAAGSARFUN   NMGRPPPRATTIPPGAA*   REEAPGKRYSPPGPSGA   TGPRAGFQAKGAAGSARFUN   TGRAGFQAKGAAGAAGSARFUN   TGRAGFQAKGAAGAAGAAGAAGAAGAAGAAGAAGAAGAAGAAGAAGA							KCRTAAGTRQWGGRQSQLQRC
RGPGGKSGFVKVQGPH   LGTWCLVSQFLHPiQ*KG   SGVASEGGSFKPWQLPR   MGAQKSRTEVWEPLPRE   MGAQKSRTEVWEPLPRE   GNTWMPRIPMCARDEG   GCSNPSQRR   GCSNPSQRR   GPRHADPANFSARNLSRT   GRVQPPRPQAPQLIPPGLVRPPPAPQAPQLIPPGLVRPPPRPVDRTLPGGP   AAAAOPTGGRSGSPRAGG   TGPRAGPARQAAAOPTGGRSGSPRAGG   TGPRAGPARQAAQAAAOPTGGRSGSPRAGG   TGPRAGPARQAAQAAAAOPTGGRSGSPRAGG   TGPRAGPASRAGAGSAAGSAP   NMGRPPPATTIPPGAA*   REEAPGKFRVSPPGPSGA   TRASFITIAPGGAGGGDSB   IILUADAQPDSARSPPGL   LSAAPRSRRDPPDAVGGL   AARRINITWKLPQLGAS   LSAAPRSRRDPPAVGGL   AARRINITWKLPQLGAS   GNVIQKKVIKQHIPLE   SIKDEGDLRNGWVIKTPT   VYAATATEKSEWMNHIN DLLSKSGKTPSNEHAAV   SEATVCMRCQKAKFTEV   HCRKGGFVCPSEKRI   QSSKPVRICDFCYDLLSA   TCOPARSDSVSQLKSPLJ   DDDDDDDMQ   DDDDDDMQ   DDDDDDMQ   DDDDDDMQ   DRTNIMVGYHQTNQKTD   PTRRPULDHNRLPEGSS   AVAVAGDIPAAAVQVTM   DTGFDVFSGSIAESWGQ   PSYC*DWRATMLRAPF   GETKDLROFFNIDNAYMI   GETCHLORY   GETCHLOR							
LGTWCLVSQPLHPIG*KG   SG\VASEGSPKPWQLPR    MGAQKSRTEVWEPLPRF    GNTWMPRIPMCCARDPG    GCSNPSQRR    GCSNPSQRR    GCSNPSQRR    GPRHADPANFSARNLSR    GRVQDPRQAPQLPPCL    VRRPPRRPVDKTLPGGP    AAAGPTGGRGSPRAGG    TGPRAGPGARGAGASAP    NMGRPPPATTTPPGAA*   REEAPGKRVSPPPGASAGAGAP    NMGRPPPATTTPPGAA*   REEAPGKRVSPPPGASAGAGAP    NMGRPPPATTTPPGAA*   REEAPGKRVSPPPGASAGAGAP    NMGRPPPATTTPPGAA*   REEAPGKRVSPPPGASAGAGAP    NMGRPPPATTTPPGAA*   REEAPGKRVSPPPGASAGAGASAP    NMGRPPPATTTPPGAA*   REEAPGKRVSPPPGASAGAGASAP    NMGRPPATTTPPGAA*   REEAPGKRVSPPPGASAGAGASAP    NMGRPPATTTPPGAA*   REEAPGKRVSPPGASAGAGASAP    NMGRPPATTTPPGAA*   REEAPGKRVSPPGASAGAGASAP   NMGRPPATTTPPGAA*   REEAPGKRVSPPGASAGAGAGASAP   NMGRPPATTTPPGAA*   REEAPGKRVSPPGASAGAGAGAGABAGAGAGAGAGAGAGAGAGAGAGAGAGA	22694	53062	Α	22822	1	410	MGKMSPGHVRDFHGRPSHHRP
SGVASEGGSFKPWQLPR   MGAQKSRTEVWEPLPRF   GNTWMPRIPMCCARDFG   GCSNPSQRR   GCSNPSQRR   GCSNPSQRR   GGSNPSQRR   GFRHADPANFSARNLSRF   GRVQDPRQAPQLPPGL   VRRPPPRRPVDRTLPGGP   AAAACPTGGRGSPRAGG   TGPRAGPGARGAAGSAP   NMGRPPPATTTPPGAA*   REEAPGKRVSPPGPSGA   TRASPTPTAAGGGGOSS   IILLVADAGPDSARSPPGL   LSAAPRSRRDPDA VGGLA   AARRINTTWKLPQLGAS   IILLVADAGPDSARSPPGL   LSAAPRSRRDPDA VGGLA   AARRINTTWKLPQLGAS   GKMVDCLANSEANTRRISI   FGAAGQPLTIPGRVLIGEG   KLVCRKKPKARGFELRRD   GRIVQKKKYLKGKHIPLE   SIKDEGDLRGWVIKTPT   VYAAATATEKSERWMHIN DLLSKSGKTPSNEHAAVV   SEATVCMRCQKAKFTEV   HCRKGGFVVCPSEKR   QSSKPVRICDFCYDLLSA   TCOPARSDSYSQLKSPLJ   DDDDDDDMQ   DDDDDDMQ   DDDDDDDMQ   DDDDDDMQ   DRTNIMVGYHQTNQKTD   PTRRPVLYDHNRLPEGSS   AVAVAGDHPAAVQVTM   DTGFDPVFSGSIAESWGQ   PSY***DWFAATMLRAFP   GETKDLRGFRUDNAYMI   GETKDLRGFRUDNAYMI   DTGFDPVFSGSIAESWGQ   PSY****DWFAATMLRAFP   GETKDLRGFRUDNAYMI   GETKDLRGFRUDNAYMI   GETKDLRGFRUDNAYMI   DTGFDPVFSGSIAESWGQ   PSY****DWFAATMLRAFP   GETKDLRGFRUDNAYMI   GETKDLR							RGPGGKSGFVGKVQGPHAVCS
MGAOKSRTEVWEPI.PRE							LGTWCLVSQPLHP\G*KGPMYS
GNTWMPRIPMCCARDPG			1				SG\VASEGGSPKPWQLPRGVEP
GCSNPSQRR							MGAQKSRTEVWEPLPRFQKIY
22695							GNTWMPRIPMCCARDPGGAKG
GRVOPPRQAPQLIPPGL   VRRPPPRRPVDRTLEGE    AAA/OPTGGGSSPRAGG    AAA/OPTGGGSSPRAGG    AAA/OPTGGGSSPRAGG    TGPRAGPGARGAAGSAP    NMGRPPPATTIPPGAA*   REEAPGKRVSPPGPSGA   TRASPIPTAAGGGGGDS    IILVADAG/PDSARSPPGL   LSAAPRSRCPPDAVGGL/   AARRINITWKLPQLGAS   LSAAPRSRCPPDAVGGL/   AARRINITWKLPQLGAS   EVACOPLIPPGRVIGE     KMVDCLANSEANTRRISI     FGAAGQPLTIPGRVLIGE     KLVGKKPKARGPELRID     GRIVNGKKYKIKGHIPLE     SIKDEGDLRIGWVIKTPT     VYAATATEKSEWMHIN     DLLSKSGKTPSNEHAAV     SEATVCMRCQKAKFTEV     HCRKGGFVCPGSEKR     QSSKPVRICDFCYDLLSA     TCQPARSDSYSQSLKSPL     DDDDDDDMQ     22698   \$3066   A   22826   1219   2370     22698   53067   A   22827   164   601   DRTNIMVGYHQTNQKTD     PTIRRPULDHNRLPEGS     AVAVAGDIPAAVQVTM     DTGFDPVFSGSIAESWGQ     PSYC*DWEATMLRAFP     GETKDLRGFRUDNAYMI     GETKD							GCSNPSQRR
VRRPPPRRPVDRTLPGGG   AAA(GPTGGRGSPRAGG   AAA(GPTGGRGSPRAGG   TGPRAGCARGAAGAAGAP    NMGRPPPATTTPFCGAA-   REEAPGKFRVSPPGPSGA     TRASPTPTAAGGGGGDS    II.LVADAGPDSARSPPGL   LSAAPRSRRDPDAVGGL/   AARRLNTTWKLPQLGAS     A 22824   271   426     KMVDCLANSEANTRRISI     FGAAGQPLTIPGRVLIGEC     KLVCRKKPKARGFERND     GNIVIQKKVIKQHIPLE!     SIKDEGDLRNGWVIKTP     VYAATATEKSEWMHIN     DLLSKSGKTPSNEHAAVV     SEATVCMRCQKAKFTEV     HCRKCGFVVCGPCSEKRI     QSSKFVRICDFCYDLLSA     TCQPARSDSYSQSLKSPLI     DDDDDDDMQ     22699   53067   A 22826   1219   2370     DRTNIMVGYHQTNQKTD     PTRRPVLOPINRLPEGS     AVAYAGDIFBAAVQVTM     DTGFPPVFGSIAESWGG     PSYC*DWEAATMLRAFPI     GETKDLRQFFNIDNAYMI     GROUNT     G	22695	53063	A	22823	264	979	GPRHADPANFSARNLSRP*DAP
VRRPPPRRPVDRTLPGGG   AAA(GPTGGRGSPRAGG   AAA(GPTGGRGSPRAGG   TGPRAGCARGAAGAAGAP    NMGRPPPATTTPFCGAA-   REEAPGKFRVSPPGPSGA     TRASPTPTAAGGGGGDS    II.LVADAGPDSARSPPGL   LSAAPRSRRDPDAVGGL/   AARRLNTTWKLPQLGAS     A 22824   271   426     KMVDCLANSEANTRRISI     FGAAGQPLTIPGRVLIGEC     KLVCRKKPKARGFERND     GNIVIQKKVIKQHIPLE!     SIKDEGDLRNGWVIKTP     VYAATATEKSEWMHIN     DLLSKSGKTPSNEHAAVV     SEATVCMRCQKAKFTEV     HCRKCGFVVCGPCSEKRI     QSSKFVRICDFCYDLLSA     TCQPARSDSYSQSLKSPLI     DDDDDDDMQ     22699   53067   A 22826   1219   2370     DRTNIMVGYHQTNQKTD     PTRRPVLOPINRLPEGS     AVAYAGDIFBAAVQVTM     DTGFPPVFGSIAESWGG     PSYC*DWEAATMLRAFPI     GETKDLRQFFNIDNAYMI     GROUNT     G			l				GRVQDPRPQAPQL\PPGLGAPSR
AAA/ADPTGGRGSPRAGG:   TGPRAGPGARGAAGSAP    NMGRPPPATTITPPGAA*     REEAPGKRYSPPGPSGA     TRASPTPTAGGGGGDS    ILLVADAGPDSARSPPGI     LSAAPRRRDPDA VGGL     AARRINTTWKLPQLGAS     A 22824   271   426     Z2697   \$3065   A 22825   \$46   1296   KMVDCLANSEANTRRISI     FGAAGQPLTIPGRVLIGGE     KLVCKKPKARQFFLRND     GNIVIGKKYIKQHIPLE!     SIKDEGDLRNGWVIKTPT     VYAATATEKSEWMNIN     DLLSKSGKTPSNEHAAVV     SEATVCMRQKAKFTPV     HCRKGGFVVGPCSEKRI     QSSKPVRICDFCYDLLSA     TCQPARSDSYSQLKSPLJ     DDDDDDDMQ     22698   \$3066   A 22826   1219   2370     DRTNIMVGYHQTNQKTD     PTRRPVLVDHNRLPEGS     AVAVAGDHPAAVQVTM     DTGFPPVFSGSIAESWGQ     PSY***DWEATMLRAFPI     GETKDLRGFPNUDNAYMI     GETKDLRGFPNUDNAYMI     GETKDLRGFPNUDNAYMI     GETKDLRGFPNUDNAYMI     GETKDLRGFPNUDNAYMI     TGRPAGAGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGCS     TASPTCHAGGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGDS     TASPTCHAGGGGGS     TASPTCHAGGGGS     TASPTCHAGGGGGS     TASPTCHAGGGGGS     TASPTCHAGGGGS     TASPTCHAGGGGS     TASPTCHAGGGGS     TASPTCHAGGGGS     TASPTCHAGGGGS			ı			Ì	VRRPPPRRPVDRTLPGGPRPPA
TGPRAGPCARGA/GSAP  NMGRPPPPATTIPPGAA* REEAPGKFRVSPPGPSGA TRASPIPTAAGGGGGDS II.LUVADAGVPSARSPPGI LSAAPRSRDPPDA/GGL/ AARRINTTWKLPQLGAS   LSAAPRSRDPPDA/GGL/ AARRINTTWKLPQLGAS   A 22824 271 426							AAA\GPTGGRGS\PRAGGTAAA
NMGRPPPATTTPPGAA   REEAPGKFRVSPPPGAA   REEAPGKFRVSPPPGAA   REEAPGKFRVSPPPGAA   TRASFITIAAGGGGDS   II.LVADAGPDSARSPPGL   LSAARSRRDPDA VGGLL   AARRINTTWKLPQLGAS   A 22824   271   426     ARRINTTWKLPQLGAS   A 22825   546   1296   KMVDCLANSEANTRRISI   FGAAGQPLTIPGRVLIGE   KLVCRKPKARGFERND   GNIVIQKKKYIKQHIPLE   SIKDEGDLRRGWVIKTPT   VYAATATEKSEKWMNIN DLLSKSGKTPSNEHAAVV   SEATVCMRCQKA KFTEV   HRKGGFVVCGPCSEKRI   QSSKPVRICDFCYDLLSAA   TCQPARSDSYSQLKSPLI   DDDDDDDMQ   DDDDDDMQ   22699   53067   A 22827   164   601   DRTNIMVGYHQTNQKTD   PTRRPVLVDHNRLPEGSF   AVAVAGDIFBAAVQVTM   DTGFPPVFSGSIAESWGQ   PSYC*DWEAATMLRAFPI   GETKDLRQFPNIDNAYMI							
REEAPGKERVSPPGPSGA			1				
TRASPIPIAAGGGGGDS   IILVADAGPDARSPPRI   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPPO   LESA PRISREPI   LESA P			1				
II.LVADAGIPDRARSPPGL			1				
LSA APRSRROPDA VGGLA			Į.		]		
AARRINTTWKLPQLGAS   22826   271   426   KMVDCLANSEANTRRISI   FGAAGQPLTIPGRVLIGEG   KLVCRKPKARAGPERNID   GNIVIQKKYIKQHIPLE   SIKDEGDLRNGWVIKTPT   VYAATATEKSEWMYHIN DLLSKSGKTPSNEHAAVV   SEATVCMRCQCAK FTPV   HCRKCGFVVCGPCSEKRI   QSSKPVRICDFCYDLLSA   TCQPARSDSYSQLKSPLI   DDDDDDDMQ   22699   53067   A   22827   164   601   DRTNIMVGYHQTNQKTD   PTRRPVLYDHNNRLPEGSF   AVAVAGDIFBAAVQVTM   DTGFPPVPSGSIAESWOG   PSYC*DWEAATMLRAFPI   GETKDLRQFPNIDNAYMI			1		ļ	İ	
22696   \$3064   A   22824   271   426	1	j	1				
22697	22606	52064	ļ.	22824	271	426	AARRENTTWKEPQEGASTE
FGAAGQPLTIPGRVLIGEC   KLVRKKPKARGFTLAGE   KLVRKKPKARGFTLAGE   KLVRKKPKARGFTLAGE   KLVRKKPKARGFTLAGE   SIKDEGDLRNGWVIKTPT							MANDOLANGEANT DRIGIVENO
KLICRKKPKARQFFLRND   GNIVIQKKKYIKQHIPLE  SIKDEGDLRNGWVIKTPT	22097	33063	^	22023	346	1290	
GNIVIOKKY/IKQHIPILE:   SIKDEGDLRNGWVIKTPT   VYAATATEKSEWMNHIN   DLLSKSGKTPSNEHAAVV   SEATVCMRCQCAKFTPV-   HCRKCGFVVCGPCSEKRI   QSSKPVRICDFCYDLL5A-   TCOPARSDSYSQLKSPLI   DDDDDDDDMQ   22698   53066   A   22826   1219   2370     22698   53067   A   22827   164   601   DRTNIMVGYHQTNQKTD   PTRRPVLOMNRLPEGSE   AVAYAGDHFAAVQVTM   DTGFPPVFSGSLAESWGQ   PSYC*DWEAATMLRAFPI   GETKDLROFFNIDNAYMI		i	1				
SIKDEĞDLRAĞW'İKTPT VYAATATEKSEWMNHIN DLLSKSĞKTESNEHAAV\ SEATVCMRCQKAKFTEV\ HÇRKĞĞFVVÇPĞSEKRİ QSSKPVRICDFCYDLLSA TCOPARSDSVSQSLKSPLI DDDDDDMQ		ł	1				
VYAATATEKSEWMNIND			1				
DLLSKSGKTPSNEHAAVV   SEATVCMRQKAKFTPV  HCRKGGFVVGPCSEKRI			1				
SEATVCMRCQKAKFTPV  HCRKCGFVVCGPCSEKRI			1				
HCRKGGFVVČQPCSEKRI			1				
QSSKPVRICDFCYDLLSAG TCQPARSDSYSQSLKSPLI DDDDDDDMQ   22699   53067   A   22826   1219   2370   DRTNIMVGYHQTNQKTD   PTRRPVLVDHNRLPEGSF   AVAVAGDHPAAVQVTM   DTGFPPVFSGSIAESWGQ   PSYC*DWEAATMLRAFPI   GETKDLRQFPNIDNAYMI			1				
TCQPARSDSYSQSLKSPL1		l	1				
DDDDDDDMQ		l	1				QSSKPVRICDFCYDLLSAGDMA
22698   53066   A   22826   1219   2370     DRTNIMVGYHQTNQKTD			1				TCQPARSDSYSQSLKSPLNDMS
22699 53067 A 22827 164 601 DRTNIMVGYHQTNQKTD PTRRPULVOHNRLPEGSI AVAYAGDIFBAAVQVTM DTGFDPVFSGSIAESWGQ PSYC*DWEAATMLRAFPI GETKDLRQFPNIDNAYMI						l	DDDDDDDMQ
PTRRPVLVDHNRLPEGSF AVAVAGDHPAAVQVTM DTGFDPVFSGSIAESWGQ PSYC*DWEAATMLRAFPI GETKDLRQFPNIDNAYM							
AVAVAGDHPAAVQVTM DTGFPPVFSGSIAESWOG PSVC*DWEAATMLRAFPI GETKDLRQFPNIDNAYMI	22699	53067	A	22827	164	601	DRTNIMVGYHQTNQKTDTGKT
DTGFPPVFSGSIAESWOG PSYC*DWEAATM.RAFPI GETKDLRQFPNIDNAYMI			1				PTRRPVLVDH\NRLPEGSRGRL
PSYC*DWEAATMLRAFPI GETKDLRQFPNIDNAYMI	1		1		1	l	AVAVAGDHPAAVQVTMTLVN
GETKDLRQFPNIDNAYME			1				DTGFDPVFSGSIAESWGQHSGT
		l	1		1	1	PSYC*DWEAATMLRAFPLAKK
NRADAVI HDTPNII YEME	1	l	1				GETKDLRQFPNIDNAYMELGT
I I I I I I I I I I I I I I I I I I I		l	1				NRADAVLHDTPNILYFMQF
22700 53068 A 22828 2 91	22700	53068	Α	22828	2	91	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
i	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22701	53069	Α	22829	1	1188	MGAEPSLLTSLGTLWSGERFOC
		i				MRAKIWERSRKGRRFSTDVISP
1						TKDSFAGPFQPISDTLVQHQRIY
ŀ	İ	ĺ				GLKVCVLPDPCKIRILKPNPQC
İ		ŀ				VGKWEEEVAENEIQVFGGERF
	İ		l			DINKYSKSPPPPTPQPLYLTAGF
ŀ		ı				ASCRNHRKSEELREGKGESPGK
		l				SKPFHLQANFTVALVTTLQGDK
						HSQEAFIRPVAKAKTTCTYLSN
		1				VCEMKDFKRINCPDPFHKMAP
	1	1				KAKKEA\PAPP\KAEAKAKGFK
1						GPRRQVLK\GVHNHKKRKRKK
		1				ERSRTFTHLSRRPGRPLRI\RRQP
	l	1				KYPSGRAAPRRNKLD\HYAIIK
		i	1			FPLT\TESA\MKK1EDNNTL\VFI\
		l				VELLKATTKNQIKQGC*RSLYD
1		1				I\DVGQGSTPWISAWMGEKKGI
						CSQLAPDLPIAFGMFANKIWGFI
22702	53070	A	22830	1413	2197	RVEWRWYNVLPCLDVHSGPVD
1						RRSHVMTAAEKPPFCPAWGQA
İ		1				VSQQSCALEPRTGEEGGKGIQG
			ŀ			PASGLFPTGEEISVVPNRWLDA
1						SLSPVILGVQG\GSQCLSCGVGQ
ŀ						EPTLTLEVRLGASSLGTQPQML   SLLKPGSPQPWCCGTP*QDSVD
ľ		l				GSFASSLRILAKALLCGPGFCPS
l						LTLPSFCRKPVTIME\LYLGAKE
ŀ		l				S\KSFTFYRRDMG\LT\SSFESAA\
ŀ		l				YPGWFLCTEPE\ADOPVRLTOL
	ŀ	l				PENGGWNAPITDFYFQQ\CD
22703	53071	A	22831	100	177	KRDHF*SRLQQVSRSVRWSSSA
22/03	33071	ľ	22031	100	l'''	TDSR
22704	53072	A	22832	610	1080	1DSK
22704	53072	A	22833	1169	1308	
22703	22013	<u></u>	22033	1107	1300	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop eodon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequenee	deletion, \=possible nucleotide insertion)
				sequenec		
22706	53074	Α	22834	l	1889	ACRRRAGTWTPPRPRGSGSRA
						MAGNCGARGALSAHTLLFDLP
		l				PALLGELCAVLDSCDGALGWR
			1			GLAERLSSSWLDVRHIEKYVDQ
						GKSGTRELLWSWAQKNKTIGD
	İ					LLQVLQEMGHRRAIHLITNYGA
	l	ĺ				VLSPSEKSYQEGGFPNILFKETA
						NVTVDNVLIPEHNEKGVLLKSS
						ISFQNIIE\GTRNFHKDFLIGERG
		i				E/IFGGYQEGKIQTLTYAVKLFK
						QEKKMQCKKHWKRFLSELEVL
						LLFHHPNILELAAYFTET\EKFC
					,	LIYPYMRNGTLFDRLQCVGDT
						APLPWHIRIGILIGISKTIHYLHN
						VQPCSVICGSISSANILL\DDQFH
						PKLTDFSMAHFRSP\LEHQSCCI
						NMPSS\SRKHLWY\MPEEYIRQ
						GKLSIKTDVYSFGIVIMEVLTGC
						RVVLDDPKHIQLRDLLRELMEK
						RGLDSCLSFLDKKVPPCPRNFS
						AKLFCLAGRCAATRAKLRPSM
						DE\V*RALESTQASLYF\AGDPP
						TSLKALRGP\SPLF\LEKVT\SIPV
						\EDDE\KPNNNLLPS**KALRMK
						RWPQK\TPF\ECSQP\EVWFLSLE
						QKAREARENEEACNMPSSS\CE
		l				ESWFPKY\IVPS\QDLRPL*GK*K
						ILSSQA\SGHSC\RTKPVESSCSS
						KFSWDEYEQYKKE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Uaknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22707	53075	A	22835	1	1560	RWERGGKRFRAADPMSALRRS GYGPSDGPSYGRYYGPGGGDV PVHPPPPLYPLRPEPPOPPISWR
1		1				VRGGPAETTWLGEGGGGDG
		ı				YYPSGGAWPEPGRAGGSHOEO
	1	ı				PPYPSYNSNYWNSTARSRAPYP
		l				STYPVRPE\LQGQSLNSYTNGA
	1	l				YGPTYPPGPGANTASYSGAYY
		1		1		APGYTQTSYSTEVPSTYRSSGN
1	i	l				SPTPVSRGIYPOODCOTEAPPLR
1	1	1		ļ		GQVPGYPPSQNPGMTLPHYPY
		l				GDGNRSVPQSGPTVRPQEDAW
		1				ASPGAYGMGGRYPWPSSAPSA
	1	ı				PPGNLYMTESTSPWPSSGSPQSF
		l				PSPPVQQPKDSSYPYSQSDQSM
		l				NRHNFP/CASVHOYESLGDO*T
		l				MDDSRILLDSPSPSIVPEPSACM
	İ	ı				GNAHPVTHP\NNODOSSSLPEE
		1				CVPSDESTPPSIKKIIHVLEKVO
		ı	1			YLEQEVEEFVGKKTDKAYWLL
		1	1			EEMLTKELLELDSVETGGQDSV
1		l				RQARKEAVCKIQAILEKLEKKR
1		1				IMKGFITK WKPVTNLTKEHLIW
		ı		1		LITLFLKCLLMTRSNTFQLFL
22708	53076	A	22836	8	207	TPVPRALRSPAGLEGSHLSSIST
		1				MMPAVMSVGKVTENGGSPQGI
						KSPSKPPGPNRIG*RNQVSVHT
22709	53077	Α	22837	459	5520	RQVRPAGLEGSDLSSINTMMSA
	1	ı				VMSVGKVTENGGSPQGIKSPSK
1		l		l	]	PPGPNRIGRRNQETKEEKSSYN
i				ŀ		CPLCEKICTTQHQLTMHIRQHN
		l				TDTGGADHSCSICGKSLSSASSL
		ı				DRHMLVHSGERPYKCTVCGQS
		l				FTTNGNMHRHMKIHEKDPTSA
		1				TATAPGRGDFASAKSSKRKLSH
	1	1				DAESEREDPAPAKKMVEDGQS
		1				GDLEKKADEVFHCPVCFKEFV
		1				CKYGLETHMETHSDNPLRCDIC
22710	62070	١.	22838	164	424	CVTFRTHRGLLRHNAL
22710	53078	A	22839	164	242	
22711	53080	A	22840	187	325	
22713	53081	A	22841	1	130	NCPMCWPGOSPRVPST*GRRNP
[227.13	23001	ľ	22041	ľ	1.50	PPKRRSVSAES*RHWPQSPSP
22714	53082	A	22842	614	774	OEPMRMCTMLANSSPSSSOATP
-2//		ľ.		1	l'''	PGL*SWNQPSLPARTLSRHEWT
			l		1	SCLGSSANL
22715	53083	A	22843	396	560	LFPHDGIKHKGKENP/PKGPPPP
	1	1		I		THPOTPPRPADAPLVATKSTLD
						OAWALLAVATS
22716	53084	В	22844	124	205	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence	1	09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22717	53085	-	22845	1	2355	MVGSRRISGVIPLPFTRRAGIFS
22/1/	23083	A	22843	1	2333	
		1				DPRANMQPLWAPVPPATFVPTS
	İ					LPPVPTARPPTSASCHCQQVPC
		i				AGLEVALPGNRTICWLSLWPS
	İ	1				WQLDEDAQSPALESSSIRQFPT
	l	1				APRSGYPGTLCPSFLFGPRPRTC
	ŀ					EYHSKPVLCAPSSRASPSSATTI
		1			1	STRTLWGPGAGSHPFGVHNTR
						LSPDLCPGKIVLRALKESGAGM
		1				PEQHKDPRVQENPDDQRTVPE
		l		i	į.	VTGDARSAFWPLRDNGGPSPFV
						PRPGPLQTDLHAQSSEIRYNHTS
					ŀ	QTSWTSSSTKRNAISSSYSSTGG
		1				LPGLKQRRGPASSRCQLTLSYS
						KTVSEDRPQAVSLGHTRCEKG
			İ			ADTAPGOTIAPTASRPHINTSLH
		1			1	VEDKAISDCRPSRPSHTLSSLAT
İ		l				GASGGPPVSKAPTMDAQQDRP
		1			ļ	KSQDCLGLVAPLASAAEVPAT
		1				APVSGKKHRPPGPLFSSSDPLPA
		1				NSSHSRDSAQVTSMIPAPFTAA
		1			1	SRDAGMRRTRSAPAAAAAAPP
		1				PSTLNPTSGSLLN\EWMEAPHIS
						WPOPOLOHVPRGO/NVRYNOR
		1				SQTSRTRS/CPQTKCQLQLPQLY
l						GRPPGSKAEEG\PASSHCQLALS
		1				SSNTVSEDGPQAVSSGHRCENK
		l				AGTAPGQTLAPRGGSPRSQASR
1		l				PHINTALHVEDKAISDCRPSRPS
1		1				
						HTLSSLATGASGGPPVSKAPTM
1						DAQQDRPKSQDSLGLLAPLASA
		ļ		l		AEVPSTAPVSGKKHRPPGPLFSS
22718	53086	Α	22846	1	1200	MPEQDKDPRVQENPDDQRRVP
				1		EVTGDARSAFRPLRDNGGLS/PF
1						FVPGPGPLQTDLHTQR\LGYRV
1		1		l		TVEDLDREKEAAFQRINSALQV
1				l	i	EDKAISDCRLSRPSHTLSSLATG
			ł	l		TSGLPAVSKAPSMDAQQERHK
l				l		SQDCLGLLAPLAPAAEVPSTAP
		l		l		MSGKKHRPPGPLFSSSDPLPATS
1						SHSQDSAQVTSLIPAPFPAASM
				l		DAGMRRTSPGTSAPAAAAAPPP
1						STLNPTLGSLLEWIEALHISGPQ
1	1	ı	1	I		PQLQQVPRGQNQRSQTSRTSSC
	1	1	1	1	1	PKRK/CHLQLLQLYGRPPGTKA
						EEG\PASSHCQLTLSSSNTVSED
1	1		l	1		GPQAVSSGHTHCEKTADTAPG
1	1	1	i			QTLAPRGGFPRSQASRPCRHKF
		1	l	l		PLLPRRRGEPLMLPPPLELGYR
		1				VTAEDLDREKEVAFQRIKSALQ
1	1	1		1		VEDKAI
					l	VEDRAI

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22719	53087	Α	22847	2	1762	YHSKPVLCAPSSRASPSSATTLS
						TRTLWGPGAGSHPFGVHNTRL
		1				SPDLCPGKIVLRALKESGAGMP
ļ	į.					EQHKDPRVQENPDDQRTVPEV
						TGDARSAFWPLRDNGGPSPFVP
		ŀ				RPGPLQTDLHAQSSEIRYNHTS
ł						QTSWTSSSTKRNAISSSYSSTGG
		ŀ				LPGLKQRRGPASSRCQLTLSYS
	1	1				KTVSEDRPQAVSLGHTRCEKG
		ŀ				ADTAPGQTIAPTASRPHINTSLH
		1				VEDKAISDCRPSRPSHTLSSLAT
l		ŀ				GASGGPPVSKAPTMDAQQDRP
		[				KSQDCLGLVAPLASAAEVPAT
l		l				APVSGKKHRPPGPLFSSSDPLPA
	ł	1				NSSHSRDSAQVTSMIPAPFTAA
		1				SRDAGMRRTRSAPAAAAAAPP
		ł				PSTLNPTSGSLLN\EWMEAPHIS
	İ					WPQPQLQHVPRGQ/NVRYNQR
	1	l				SQTSRTRS/CPQTKCQLQLPQLY
ļ				1		GRPPGSKAEEG\PASSHCQLALS
	İ	1		ļ		SSNTVSEDGPQAVSSGHRCENK
	ł					AGTAPGQTLAPRGGSPRSQASR
ŀ		1				PHINTALHVEDKAISDCRPSRPS
				1		HTLSSLATGASGGPPVSKAPTM
			1		}	DAQQDRPKSQDSLGLLAPLASA
					f	AEVPSTAPVSGKKHRPPGPLFSS
1						SDPLPATSYHSRDTAQ

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22720	53088	A	22848	292	2053	GSVAWSPFSRSSLCLWDLPV APTLWGPGAGSHPFGVHNTRL SPDLCPGKIVLRALKESGAGMP EQDKDLRVQENPDDQRRVPEV TGDARSAFLAPAGQWSPLSLC AQSRASADIPPCFEVRQI*PDIP HLDELMIPOKCHLQLLQLCG RLAGAKVEEGASGAKEQGRDA "AGQGTPESQQNPDAGREYEV RGPLQTDLHAQSSEIRYNGTS OTSWTSSTKRNAISSSYSSTGG LIGLKQRRGPASSRCQLTLSYS KTVSEDRPQAVSSGHTRCEKA ADTAPGQTLAPRGGSPRSQGSR PRRRKIALLPHR*GELQVEDKAI LDCRFSWPSHTLSSLATGASGE PPVSKAPTMDAQQDRPKSQDC LLGLVAPLASAAGIPSTAPVSGK KHRPPGFLFSSSDPLPATSSHISS DSAQVTSLIPAAFTAASMDVG
						LNPTSGSLLN\EWMEALHISGPQ PQLQQVPRGQ/NVRYNQRSQTS RTRS/CPQTKCQLELPQLYGRPP
						GTKGCRHEKNVLCSKLFEGFGF IFVGFFFFAYMGILQLLII

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22721	53089	A	22849	146	1789	RQPSTSGSSRSCPCARRSWWAR
		l				TARSGWWWELLGARRSPOTLH
		l				CPQATLITLPCPRPSSTTSGSAM
		ŀ				T*RGPWRSPGCTTSFCPTSR/PV
		1				ERNIDQTDLHAQRSEIRYNQTS
		1				QISWTSSCTNRNAISSSYSSTGG
		i				LPGLKRRRGPASSHCQLTLSSS
		1		1		KTVNSDKFDANDPILKDQTQE
l		1				WSGSATFTSDGKIRLFYTDYSG
						KHYGKQSLTTAQVNVSKSDDT
ĺ						LKINGVEDHKTIFDGDGKTYQN
İ						VQQFIDEGNYTSGDNHTLRDPH
		l				YVEDKGHKYLVFEANTGTENG
						YQGEESLFNKAYYGGGTNFFR
		1				KESQKLQQSAKKRDAELANGA
1		1				LGIIELNNDYTLKKVMKPLITR
		ı				GSSRPSIREDENGRSQKPVHPEG
						DMTMNIKKIVNQATVLTFTTAL
		1				LAGGATQAFAKENNQKAYKET
		1				YGVSHITRHDMLQIPKQQQNEK
		1				YQVPQFDQSTIKNIESAKGLDV
						WDSWPLQNADGTVAEYNGYH
		1		1		VVFALAGSPKDADDTSIYMFY
		İ			1	QKVGDNSIDSWKNAGRVFKDS
		L				DKFDANDPILKDQTQEWRGEA
22722	53090	A	22850	500	1034	

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide Jocation of first	Nucleotide location of last	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
NO.	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22723	53091	Α	22851	1	2396	MDKYLLHGPPRPPPAIASRCLA
						RVPGCAYSSIMVALSLVPLCAH
						MPTPSGLEVALPGNRTICWLSL
		1				WPSWQLDEDAQSPALESSSIRQ
		l				FPTAPRSGISPLCAPHSCLGQGF
	ľ	l				GPASHPTSTQTLDISGVKTTPRI
		l				FSVILQDLCSILGDSCLQFHLHR
	į.	l	Į.			VPQQVSAVLLRVSGASPSPATT
		•				LSTRTLWGPGAGSHPFGVHNT
	ł					RLSPDLCPGKIVLRALKESRAG
		l				MPEQDKDPRVQENPDDQRTVF
						EVTGDARSAFWPLRDNGGPSP.
	1		İ			VPRPGPLQTDLHAQSSEIRYNH
						TSQTSWTSSNTKRNAISSSYSST
						GGLPGLKQRRGPASSRCQLTLS
	1	1				YSKTVSEDRPQAVSSGHTRCE
		i				GADTSPGQTIAPTGGSPRSQDS
						RPRRKIPLLPRRRGEPLMLPPF
	ļ	l		1		LELGYRVTAEDLHLEKETAFQI
				1		INSALHVEDKAIPDCRPSQPSH
	l					LSSLATGASGGPPVSKAHTMD
						AQQHRPKSQDCLGLLAPLASA
						AEVPSTAPVSGKKHRPPGPLFS
		1				SDPLPATSYHSRDSAQVTSLIPA
		l		l		PFTAASSDAGMRRTRPGTSAPA
	l			ŀ		AAAAAPPPSTLNPTSGSLLN\EV
						MEAPHIS WPQPQLQHV SRGQ/N
						VRYNQRSQTSRTRS/CPPRKCQ
				1		QLPQLYGRPPATKAEEG\PASSI
	l .			1		CQLAHSSSNTVSEDGPQAVSSC
				1		HTRCEKKAGAS/RWASRF*STH
		_				YGCTAGQTQVPRLPGPTGPPSI
22724	53092	A	22852	1	3392	MDKYLLHDPPRPPPAIASRCLA
	i					RVPGCAYSSIMVALSLVPLCAF
		1				MPTPSGLEVALPGNRTICWLSL
		1				WPSWQLDEDAQSPALESSSIRC
		ı				FPTAPRSGYPGTLCPSFLFGPRP
						RTCEYHSKPVLCAPSSRASPSS/
		l				TTLSTRTLWGPGAGSHPFGVHI
		1				TRLSPDLCPGKIVLRALKESGA
		1				GMPEQDKDPRVQENPDDQRT\
		1				PEVTGDARSAFWPLRDNGGPSI
			l			FVPRPGPLQTDLHAQSSEIRYN HTSOTSWTSSST
	1	1	l			
22726	52002	_	22062	40	400	
22725	53093	A	22853	40	409	WKMRHETRYNVLVSRGVGCH
22725	53093	A	22853	40	409	WKMRHETRYNVLVSRGVGCH RLFFLNGVFEHLGRAFVQRRK
22725	53093	A	22853	40	409	WKMRHETRYN VLVSRGVGCH RLFFLNGVFEHLGRAFVQRRKV DTEHQLLRAA VPANQEHRGLT
22725	53093	A	22853	40	409	WKMRHETRYN VLVSRGVGCH RLFFLNGVFEHLGRAF VQRRK V DTEHQLLRAA VPANQEHRGLT QSQHVDGHSLLLLTLPNSFLCQ
22725	53093	A	22853	40	409	WKMRHETRYNVLVSRGVGCH RLFFLNGVFEHLGRAFVQRRKV DTEHQLLRAAVPANQEHRGLT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ 1D NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22727	53095	A	22858		356	MAGPVKDREAFQRLNFVYQVS LRQGPHGDGARRPRVTAPLPQ AAHCVLAQDPENQALARFYLWH TERTIAKRLVLRRDPSVKEELL CRGCSSLLVPGLTCTPAPRDRC RGQRWTGTDLP
22728	53096	A	22859	2	270	
22729	53097	A	22860	2	364	IFSALPAPRRKVGLNLAPVTEPR DQPWAMIIDVFISRYSGSRGQPR QTLTIKGELKVLMEKELPG\FL QSGKDKDAVDKL\LKDPGRPM GDAQGGTFSEVHPCSVAAIT\SA CHK\YFE\KAGLK
22730	53098	Α	22861	282	469	DCGSELQAPPPPSLPGPLCPSIM PA/PPYASPPMMPMMGPPFRGM MPVGLAPGPMLLQYNTCS
22731	53099	А	22862	7	522	SVWWNSNGLQSMMPKFYCDY COTYULTHDSPSVRKTHCSGRK HKENVKDYYQKWMEEQAQSLI DKTTAAFQQGKIPPTPFSAPPA GAMIPPPPSLPGPPRFGMMPAP HMGGPPMMPMMGPPPPGMMP VGPAFGMRPPMGGHMPMMPG PPMMRPPARPMWPTRPGMTR PDR
22732	53100	Α	22863	3	455	WFLIHDRTHTGERPYKCNECY KAFHRSTCHAHKRTHTGEKP YECNQ*GKAFNSSHSPQIHERT HTGEKPYECKECGKAFKGPSSV RRHERTHSTKKPYECKHCGKA LSYLTSFINHLGMHTGEISHTC KICGKAFYSPSVJQTHE*THT
22733	53101	Α	22864	1	1665	
22734	53102	Α	22865	1	2650	

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		<u>.                                    </u>		1.00	laas.	Language outpopping of the
22735	53103	Α	22866	107	2384	MFGVHGTFLGIESRPPLLSLSPS
						LLYTCEMFQDPVACEDVAVNF
	İ					TQEEWALLDISQRKLYREVML
						ETFRNLTSIGKKWSDQNIEYEY
		ŀ				QNPRRSFRSLIEKKVNEIKDDSH
						CGETFTQVPDDRLNFQEKKASP
				ł	İ	EVKSCDSFVCAEVGIGNSSFNM
	l					SIRGDTGHKAYEYQEYGPKPY
						KCQQPKNKKAFRYRPSIRTQER
	i					DHTGEKPYACKECGKTFISHSG
						IRRHMVMHSGDGPYKCKFCGK
		ŀ				AVHCLRLYLIHERTHTGEKPYE
						CKQCGKSFTYSATHRIHERTHT
		1				GEKPYECQQCGKAFHSSSSFQA
						HKRTHTGGKPYECKQCGKSFS
		1				WCHSFQIHERTHTGEKPCECSK
						CNKAFRSYRSYLRHKRSHTGEK
		1		ł		PYQCKECRKAFTYPSSLRRHER
		1				THSAKKPYE/CSFQYHERTHTG
	ļ.					EKPYECKQCGKAFRCAPHLRR
		l				HGRTH/S/AEKPYECKECGKAFR
		1		l		SASHLOIHERTOTHIRIHSGERP
		l			i	YKCKTCGKGFYSPKSFQRHEKT
l		l				HTAEKPYECKOCGKAFSSSSSF
		l		i		PYHERTHTGEKPYECKOCGKA
				ł		FRSASILQMHAGTHPEEKPYEC
		1		l		KQCGKAFRSAPHLRIHGRTHTG
1		l				EKPYECKECGKAFRSAKNLRIH
		1			*	ERTOTHVRMHSVERPYKCKIC
		l				GKGFYSAKSFOIHEKSYTGGKP
ļ		1				YECGLQLPRTAAESVDSEKLQG
1		l		l		PGKQETHVQLLSLYCLRRVLDC
22736	53104	A	22867	1	391	MKEHIERNPVTIKNVVKPSINT
22/30	33104	l^	22007	l'	391	VPLEDMKAHTGEKPYEROKCE
		l		ŀ		LNQRLRDYAFQYHESNSGVVS
ĺ		l				ADSETKSGHSGCLIKEMKNATK
1		l			i	THNGITGGIIFNANHWRETNTIY
1		l				
		l				TRIHI/GEKSCDYKKCGKAFNQ
		l				YSSLRRHES\HTGEKPYERQKC
		l	l	l		ELNQRLRDYAFQYHESNSGVV
		l				SADSETKSGHSGCLIKEMKNAT
		ĺ				KTHNGITGGIIFNANHWRETNTI
		_				YTRIHINKQEKYQATHVSYAAN
22737	53105	С	22868	179	289	
22738	53106	A	22869	3	206	

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22739	53107	Α	22870	I	855	MPRAEPRATLGEQEKAGLPLG
						AWRLYLLRHFRKQTELRRSGSF
			i		i	DVTGALLVAAAVASEAVGSLR
						VAEGGPNTLLLQVLR\SW\PWC
						NKELKTMEERKVKRRSPKSFSA
						HCTQVVNVQKKMSIPVSK\ST\C
						FSNPASQVNFTSRPKVKKEVM
					1	K\EKTKPQGLEEGQRHSSQLPIQ
						HSFLT\DV\$RC\$RKMERGLL\$L\
		1				LNDFHSWKTFKHFGN\ECSI\EQ
						MEHVSGELQEKLARLNLELYG
						ELEELPEDKRKTASDSNLDRLL
						SDLEELNSSIQKLHLA\DAQDVP
22740	53108	Α	22871	67	598	SIRRHCGQRCLVRIRIWPLKLPP
						RLAVREGLRCGTLRDVMPPSM
						SGVVVGPPGP\PPGPCGDTTSGP
						APGTFGQASAPVSPPRRGHGEY
		i			1	LCSSPTPVFFPGLWQPPRAGFTA
						PERSWSCRGPRWQLRAAWAGR
		ı				LSSSPVSALLTFKVIILSTGLLFG
						PLGGWKILSSLPEIFANLAT
22741	53109	Α	22872	1	372	
22742	53110	A	22873	1	3961	FALAPEAGFYSLGQQPSTRISGS
		1			İ	GCVHPVRVWPFSIEPAIGTLNV
						GESMQLEVEFEPQSVGDHSGRL
		i				IVCYDTGEKVFVSLYGAAIDMN
						IRLDKNSLTIEKTYISLANQRTIT
		l			i	IHNRSNIIAHFLWKVFATOOEE
		l				DREKYSKVTAVFSIAASSNSRF
		l	ŀ			ALYESSYNSQRSPSEWACDDLI
		l				KEEKDETDEFFEECITDPLLREH
		1				LSVLSRTFANQRRLVQGDSKLF
		l		1		FNNVFTVEPLGWSERKEMAEIR
			l			DVQKKDTKV
22743	53111	A	22874	1248	1716	WYAEQSAHEATAQLSTPMAIR
122,43	55111	n	22074	1270	17.10	RAVVMPPEORNRTKKEY/SLRI
		l				PFGLPGDRMCFFTHRNDD/PTG
						NEVTPPSLAADLGQQFGGDAT
		[				
		l	ŀ			DRQPWMIHQMLIHQGSEQLCF VEHGINMLRLGEILHLAATRHE
				l		
		1		ı		AHFLAFRQFFNHQRSCFALNLA
22744	53112	_	22875	478	663	RQDRITDF
22745	53112	A A	22876	508	884	POPURDOPROPEI PUGUCACO
22/43	22110	l^\ .	228/0	300	004	RQRHPDQRPGRSLFHGHGMCP
						TTPLQQQVDSQHNGDNNSESA
				I		P*QRRADSIHRSSLHIKKRPPSE
				1		DVSRCFLCIDLAFRQGFTFNKK
						GIRVRIPFGLPGDRMCFFTHRN
		<u> </u>				DDRPAMKLLPFAGNKV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22746	53114	A	22877	li .	506	AFCRFSILVGSAFISETDMCLLC
		1				PFPIGKAISVRLQYFKHETKQSI
l			l			CLGNOKSPLVCPFFGEOGLDSS
		1				SSLDLPSSPNPAPTLCPCVOPPR
						GPVSVSVPVSVMGSRLAPLLSA
						*LFVSVILGRMVILKNPGVLGQ
						ROAGPSPGAPGLPSPSVRAPLG
						HKCSERSPSAT
22747	53115	A	22878	621	877	RRRRRRKQRRKQPWSSPARGA
1						*PGPRRPSLRAAGTRRRRKRRA
		l				SWTRCSNTCPGRCRTCSRSNQA
		ļ				SCPSLLHLLLLLFGSFFSFY
22748	53116	Α	22879	Ī	1339	MPSYKGCGETQRHATNGTAAP
	1	l				PRRSSCRGDAIKLPHAEKKEGR
		1				PPASTSYPLRLGTRSARSSSHC
		1				WCTSWEDPPEATAGEEKRHPK
		1				SPAVCCGEGTVLTASTVTWRA
		ı				EPARGEIWARLCPRDRPGSRAA
		i				AVIWERPCCDRLIADRTGPPNSS
	1	i .			İ	EPHCRAATIPLPTWPLPEGOTLT
	1	l				LKALAKMAAPALLLLALLLPV
		l				GAWPGLPRRPCVHCCRPAWP/L
	ŀ	l				WTLCPGEKGEAGVRGRAGRSG
		l			i	KEGAA\GCPGPAGPQSARRGSV
		l				GPPGAACRRAYAAFSVGRREG
						LHSSDHFOAVPFDTELVNLDGA
		1				FDLAAGRFLCTVPGVYFLSLNV
		1				HTWNYKETYLHIMLNR/QAA/V
		1				VLYAQPSERSVMQAQSLMLLL\
	l	1	1			RRATPVWSVPEELPAVLPGODE
	ŀ		1			LGSLOLHAHHPLADGLVRHEL
	ļ	1				OPGLAGSSRTGHLVFRTDHGLR
	į.	1				OHASHGCPVGPLHVPG
22749	53117	A	22880	532	783	VGSTRGIVEGLIFILGEGGECYY
		1				PRSLFTVANFTVGKRV\KTLNC
		1				PSTDKWINKMCHIHTVEYYSDI
		1			1	KRNEIPMRATCROTLKA
22750	53118	A	22881	275	720	TPIHNCFKENKIPRNPTYKEHEG
-2.00	1	ľ		[	1	PLOGELQTTSQGNKRGYKQME
		1				EHSMLMG\GRINIMKMAILPKVI
	1	1				YRFTIPIKLPMTFFTELEKTTSKF
	1	1				IWNOKRARIAKSILSRKNKAG/
	1	1			1	AHHAT*LQTILQGYSNQNSMVL
		1			I	VPKORYRPMEOSPOK
	1			L		VENCKINEVSPOK

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \≔possible nucleotide insertion)
			1	sequence		
22751	53119	Α	22882	1	609	MGRNQSRKAENSKNQSASSTP
		1				KDHSASPATEQSWMGNDFDEL
		l				TELGFRKSVMTNFSQLKEDVRT
	1	1				HHKEAKNLEKRLDK WLTRINSI
		1			l	EETLNDLMELKTMARELRDTC
	1	1			l	TSFSSRFDQVEEGEIQTTIREYY
	İ	l		ŀ		EHLYANKLENLEEMDKFLDTY
	1			l	1	TLPRLNQEEVESLNRPITGSEIE
		1		l		AIINSLPTKKIPGPDRFTA*RRP*
		1	İ	l		MT*WS*KPWHENYVTHAQASV
	Į.	1			İ	ADSIKWKKGKYK\HHQRIL*TP
						LRK*TRKSRRNG*IPGHIHSPKT
	1	1		l		KPGRS*IPE*TNNRF*N*GNN**P
	1	L				TNQKNSRARQIHSQILPEVQRG
22752	53120	A	22883	191	564	PNARKLRTLKKDHSPD**RRKL
	i	ı				RKIK*MQ*KMIKGISQ\RSHRNT
1	1	l		l	l .	NYHQRTL*TPLHK*TGKSRRNG
	i	ı				*IPGHIHPPKSKPGRSQIPEYTNN
		l				KI*N*GSN**PINQKKSRTGRIHS
		L.				QILPEVQRGA
22753	53121	A	22884	337	589	TRRSRRNG*IPGHIHPPKTKPGR
	ľ	l		1		SRIPE*TNNRL*N*GSS**PTNQK
		i		l		KPRTRETDSRILPEVQRGAGTIP
22754	53122	ļ. —	22885	162	539	SETIPNNRKRGTPP TRKSRRNG*IPGHIHPSKTKPGR
22/54	53122	Α	22885	162	339	
1		l				S*IPEETNNKF*N*GSN**PTNQK KPRTROVHSRILPEPISHGMGKS
	İ	l				WKOSL*KPAODKAVPSLIAPIO
1		l		l		HIIGSSGQGNQARERNKGYSNR
		l				KRGSOIVSVCR
22755	53123	A	22886	329	459	NDDTFRKPHRLSPKSP*AADKQ
22/33	33123	r	22000	329	439	LOOSLRIONQCAKLRSIPIHH
22756	53124	A	22887	58	440	GLSROFSPHSVNKAAGEVPTGR
22/30	33124	<u>۱</u> ^	22007	176	1770	GPPQLSKANVARLPLSIPPLWA
1		l			1	GHLWKKGGSPSOGIIDKTLISLG
		i i				QST*GKGQLWVQLQRT*AFLPA
1		l				GSEESSGSPSTA\PNSAKGOTPS
		l			i	SSGSLTRASSSSLYLW
22757	53125	A	22888	219	332	NEGKNLKGSORERNTNYHORT
122.57	155.25	Γ.		[/	J	Q*TPLHK*TRKSRRNG
22758	53126	A	22889	1295	1435	ONOROKPHDYLNRCRK/W/HFD
-2,53	[	Ι.		1	l	KIOOPSMLKTLNKLGNDGTYL
		1			ŀ	RIIRDI
	1		L			l'ano.

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22759	53127	IA	22890	2	648	YRFSGPTIRSIONHTTHNOGRK
						WHKKGIKKPRAORYESLKGPE
						V*PSGQGVTCLLRADWLSPLLG
1						SGTY\VGPKFLRNTRFAKKHNK
				ŀ		KGLKKMQANN\AKAMSARAE
						AIKALVKPKEVKPKIPKGCS/RR
						KLRFDFAYIAHPKLWGRRCFVA
						RFCQGASGCSGPKGKAKVPKA
				i		KVKVS/PKAKAKAKAKDOTKA
				ļ		
1				i	i	QAAAPALV\PAQAPKRTQAPTK ASE
		L.	*****		370	OMYCVFNRNEDACRY/GS/AIG
22760	53128	Α	22891	3	370	
						VLAFLASAFFLVVDAYFPQISN
		l				ATDRKYLVIGDLLFSG\VLASLA
		ł		l		YQRYKAGVDDFIQNYVDPTPD
į.		1				PNTAYASYPGASVDNYQQPPFT
		_				QNAETTEGYQPPPVY
22761	53129	Α	22892	1	297	
22762	53130	Α	22893	3	713	GGGSGGDGDMESGAYGAAKA
		l		İ		GGSFNLRRFLTQPQV\VARAVV
						L\VFALIVFSCIYGEGYSNAHES
1						KQMYCVFNRNEDACRYGSA\IG
		ŀ				VLAFLA\SA\FFLVVDAYFPQISN
		1				ATDRKYL\VIGDLLF\SALW\TFL
1						WFR/VGFCFLTNQWAVTK\PKT
		1				VLVGADSVRAAITFSFFSIFSWG
		1	l			VLASLG\YQRYKAGVDDFIQNY
		1		i		VDPTPDPNTAYASYPGASVDN
				1		YQQPPF\TQNAETTEGYQPPPV
22763	53131	Α	22894	3	219	
22764	53132	A	22895	1445	1777	PVHLGSTCHQCRQKTIDTKTNC
		1		1		RNPRLLGA/CRGQFCGPCLQNR
		ı				YGEEVRDALLDPNWHCP\HCR
		1	1	1	ŀ	GICNCSFCRQRDGRCATGVLVY
		1	1			LAKYHGFGNVHAYLKSLKQEF
		1		1		EMQA
22765	53133	Α	22896	3	337	
22766	53134	Α	22897	425	1209	TMSSTQFNKGPSYQLFAEIKNQ
		1		1		LLFKCDPQKQAALCSWIEGLTR
						LSFSPDFOKSLKEGIVLCTLMN
		ł				KLGSVPKI/NRLMONWHOLENL
		1				SNFIKAMVGYDMNPMDLFEAN
		1			I	DLFESRNMTQNRIL/P/PLDHSTI
					I	SLOMGANKRSSO/VGMTAPGTP
		1			1	OHIYNTKLGAHKCDNSSMSLH
1	1		1		I	MGDTQSANHSVQGFGLGQQIY
1	1	1			I	YPKYCPO/GPAASGAPLDAGNC
1		1		1	1	
1		1				PGPGEAPEDPPCSQEEATTEAPE
		1		1	1	HAGSLHHRPVWVWGIFLYFRL
	L	11		L	L	SFFFL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22767	53135	A	22898	156	559	OHIRETPGAOVHSCGK WNPOA
						LWRTAWQLLT/DTKHTLGKPPS
						/MPHLVIYPLEISC\HTKTCTRML
			l			IASFFIITKTWKRPRCPSVDKWK
		1				NKPWCIHTGLLSSTVPPGEPGC
			İ	i		PIPELSGLLMGLOTTSSRCSVRO
						vw
22768	53136	A	22899	161	359	
22769	53137	A	22900	151	383	
22770	53138	A	22901	238	546	PQEHIMEHLEGVINKPRGRRCR
						PQELQLHYLQKCMDYDGNNLL
						DGFRTLPQPSTHV\HKEEGSEQ
						APLMSEDEL\INIIDGVLRDDDK
						NNDGY\IDYAEFAKSLQ
22771	53139	A	22902	1	819	MAAEDKLLLPQLPELFETSKQL
		l				LDDVEVVTEPAGSRIVQKKVFK
		1				GLDLLEKAAEMLSQL/DLVAEL
1			ĺ			ELPKTKNSAE\NHTANFSMAYP
						SLIAMASQRQAKRERHKQKEL
						EHRLSAIKSAVETGQADDERVR
			i			EYYLLHLQSLEEIESIDQEIKILR
						ERDSSREASTSNSSRQERPPVKP
		1				FILTRNMAQVKVFGAGYPSLAT
İ						MTVSDWYKQHWKYGALPDQG
	i					IAKATPEEFRKAAQQQEGQEEK
						EEEDDEQTLYRVREWDDWKGI
						HPRDYGSQQSMG
22772	53140	В	22903	238	410	
22773	53141	A	22904	1	638	VAELELPKTKNSAE\NHTANFS
						MAYPSLIAMASQRQAKRERHK
	į.					QKELEHRLSAIKSAVETGQADD
						ERVREYYLLHLQSLEEIESIDQE
						IKILRERDSSREASTSNSSRQERP
						PVKPFILTRNMAQVKVFGAGYP
						SLATMTVSDWYKQHWKYGAL
			l			PDQGIAKATPEEFRKAAQQQEG
	1		l	l		QEEKEEEDDEQTLYRVREWDD
			l		1	WKGIHPRDYGSQQSMG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22774	53142	A	22905	3	1083	RGNSRLRYSHEDELQLPRLPEL
		ľ.		ľ		F\ETGRQLLDEVEVAT\DPAGFR
1						\IVQEK\VFKG\LDLLE\RAAEML
						\SOSTLFGRECRFWKDLSPTHLI
						TFWGP\AFQGA\LPLKQVNPSKR
						LRIILQR\AREHFI\NYLT\QCH\C
1						YHVGQSFEL\PKT\MNNSAENH
						TANSSMAYPMLGAM\ASQRQA
İ						KIQRYKQKKELEHRLSAMKSA
i						VESGQADDERVREYYLLHLQR
						W/VDISLEEIESIDQEIKILRERDS
1		•				SREASTSNSSRQERPPAKPFILT
		l		1		RNMAQAKVFGAGYPSLP/TLW
						TVSDWYEQHRKYGSITGFRGIA
	ŀ					KSAPEGISGKAAQQQEEQEEKE
		l				EEDDEPNNFHRAREW\DDWKD
						TPSLGAYGNRQNMG
22775	53143	Α	22906	30	457	
22776	53144	Α	22907	401	1509	MWKMNCSLLQSKQMCNQLPP
•						GSTRDGSWKFRQDPRLGKNA/C
						DCDSCIASEFGVRGYPTIKLLKG
						DLAYNYRGPRTKDDIIEFAHRV
1		l				SGALIRPLPSQQMFEHMQKRHR
1		l				VFFVYVGGESPLKYVTLKEMP
1						AVLVFKDETYFVYDEYEDGDL
						SSWINRERFQNYLAMDGFLLYE
		l				LGDTDLNASSLRERKLILPEDIP
		l				LYVGKLVALAVIDEKNTSVEHT
						SELTVPTVVVLNTSNQQYFLLD
		l				RQIKNVEDMVQFINNILDGTVE
l		l				AQGGDSILQRLKRIVFDAKSTIV
		l				SIFKSSPLMGCFLFGLPLGVISIM
		l				CYGIYTADTDGGYIEERYEVSK
		1				SENENQEQIEESKEQQEPSSGGS VVPTVQEPKDVLEKKKD
22777	53145	A	22908	53	543	LPISSLRTQTLFSSPLFPQHFEHY
22111	33143	^	22900	33	343	LALVSVLPFQLGRGRTAPAKKG
		l				GEK/RKGHSAINEVVIQEYTINI
						HKRIHGVRFKKRAPRALREIPK
		1		1		FAIKEVETLSVRIDSRFNKALCA
1		1		1		KGIRNVPYRIRVOLSRKRNEDE
1		ı		1		DSPNKRYTLVTYVPVSTFKNVQ
1		l				TVHVHEN
22778	53 146	A	22909	106	491	IGKGTNELMYGEK/RKGHSAIN
1		ľ		1		EVVIOEYTINIHKRIHGVRFKKR
		l				APRALREIPKFAIKEVETLSVRI
1		l		1		DSRFNKALCAKGIRNVPYRIRV
1		1				OLSRKRNEDEDSPNKRYTLVTY
		1		1		VPVSTFKNVQTVHVHEN
22779	53147	Α	22910	1	321	· · · · · · · · · · · · · · · · · · ·
	F	1			·	

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
		<u></u>				
22780	53148	Α	22911	313	402	R*GGEKKKGRSAINEVVTREYT
						INIHKRIHGVGFKKRAPRALKEI
		l				RKFAMKEMGTPDVRIDTRLNK
	1	l				AVWAKGIRNVPYRIRVRLSRKR
						NEDEDSPNKLYTLVTYVPVTTF
						KNLQTVNVDEN
22781	53149	Α	22912	1	3801	
22782	53150	A	22913	2	535	GGGMADTVLFEFLHTEMVAEL
	İ					WAHDPDPGPGVSAGLRGEEAG
						ATKGQKMSLSVLEGMGFRVGQ
1	l	l				ALGERLPRETLAFREELDVLKF
l	l	1				LCKDLWVAVFQKQMDSLRTN
		l				HQGTYVL\QDNSFPLLLPMGSG
	ŀ	l				LQV/YLEEAPKFLAFTCGLLRSA
	ĺ					LYTLGIESVVTASVAALPVCKF
						QVVIPKS
22783	53151	Α	22914	1	201	GDLPWEINPLSSCSLLHEKDPPT
ŀ						TSGPQT/DQPKKHHTNFKSGP\T
		l				GNRTVQLTWQPLPEPLELWPK
		$\perp$				AL
22784	53152	В	22915	226	522	
22785	53153	С	22916	1	705	
22786	53154	С	22917	1	462	
22787	53155	С	22918	I	505	
22788	53156	В	22919	I	1059	
22789	53157	С	22920	I	212	LTSFLHCCLSHHYHGAQIHPRH
22790	53158	A	2292 I	1	212	FLHLICRCHCHQYCHHAYLDPC
		1				PODSHHLHWCWOPLPEPLELRP
			1			RLSD
22791	53159	В	22922	1	864	(KL3D
22791	53160	В	22922	li .	635	
22793	53161	В	22924	1	874	
22794	53162	A	22925	1	566	MPSVRTGDPIILHAHTLNPAVW
22/94	155102	^	22,23	l'	1000	LAWAVEGEGSLTLNIWICPWOI
		l				EPLRHLAYTPGSDATSLTVPTR
	ŀ	l				VISPLCVDSWLFTDPLMKTHAP
		l				ERTFLSFPAAAKROAHLCVLAS
		l				NCDKPMYVKLVKAHCAEHQIN
		1				LIKGDDNKKLGEWEGPCTIDRE
		1				GKPCKVVGCSCAVVKDY/GKE
						SQAKDVIEQYFKCKK
22795	53163	A	22926	59	602	PTAMVEEGIAAGGVMDVNTAV
22193	155105	l^	22,20		I***	QEVLKTALIHDGLARGIREAAK
		1			I	ALDKYVYOSOYCGFLOPEONC
	1	1			1	HPREEGMEFMVLAQKF*MASV
	1	1			1	LPET*EK\RQAHLCVLASNCDEP
1	1				I	MYVKLVEALCAEHQINLIKVD
	1	1	1		I	DNKKLGEWVGL\CKIDREGKPR
					1	KVVGCSCVVVKDYGKESQAK
					I	DVIEEYFKCKK
					J	PTICETIKOKK

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SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location uf first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon,/=possible nucleotide deletion, \=possible nucleotide insertion)
22796	53164	С	22927	135	293	
22797	53165	В	22928	20	325	
22798	53166	А	22929	677	1031	TPTMCLRSHPLPLALVSVILSFP QPSFSSSLINSSWSSPRSSLSSSSS RSPSSGRVNLHPPQTPSSH*SHL SSRSCCSAPLLTYHSSSLLLVCS VPFPFFPGFPGENPLFVLSGSAP
22799	53167	A	22930	1	1011	MCLRVIFTIGAVSVILSFLSLFFI LDQLQLVIPQSSLSSNLEDPY SSYYAAARRPDSASTEPPPAG KSSAEYRAHFLAVSTEAERIRA DLVLRARPLOPAHPRILUPP AGPFCRMVAHNQVA\ADNAVS TAAEFRRPEP:SPSSSSSSPAAT RAPRICPAVPAPAFGDTHFRTF RSHADVRRITRASALLDALGF YWGPLSVHGAHEPLRAEPV TFVRDSIRQRNCFFALSVKMAS GPTSIRVHFQAGRFHLDOISRES FOLTEELLEHYRGGSRADLLGA TULRQRRVRPLQELCRQRIRRPP VGJRENLLARIPLNPVLRDY\LSS FFFQI
22800	53168	В	22931	180	1466	
22801	53169	Α	22932	I	207	
22802	53170	A	22933	408	3758	CLQOGPGONQHCSHREEAPAV ACQEPFOHESPESVLPLGAPP, SEEDGRQQVGSSRLRHIMAEMI ATEREYIRCLGYVIDNYFPEME RMDLPGGLRGKHHVIFGNLEK LHDFHQQHFLRELERCQHCPLA VGRSFLRHLIGGKQFPESPSGER CTJSSHAPCFWMPHYLPEGPA CPVLRIPCEAFNWKHVECTDPV VSPPKQSPLLFQEEQFGMYVIY SYNKYQSPLLFQEEQFGMYVIY PLGPPTACPALS
22803	53171	A	22934	3	355	EAAELDAMCTATRQWSRAGQ GVTGRPCQGRLPRCGLRSTITP KPIPGGHRKKPLLGAL\GKHTLS YVERWHWEGHSGI*TRDRQNF ERRLTVPPSRQPFAPTDETRCG DDRISFCNYV

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22804	53172	Α	22935	1928	5312	CLQQGPGGNQHCSHREEAPAV
						AACQEPPGHSEPESVLPLGAPP\
	1	1				SEEDGROOVGSSRLRHIMAEMI
				1		ATEREYIRCLGYVIDNYFPEME
						RMDLPOGLRGKHHVIFGNLEK
		1				LHDFHOOHFLRELERCOHCPLA
				i		VGRSFLRHLIGOKOFPESPSGER
		1				PGLTSSHAPCPWMPHYLPEGPA
		l				CPVLRIPCEAFNWKHVECTDPV
						VSPPKQSPLLFQEEQFGMYVIY
		1				SKNKPOSDALLSSHGNAFFKDK
		1				ORELGDKMDLASY
22805	53173	A	22936	3	306	QREEGERINGERS
22806	53174	A	22937	3	460	LQRQRQHPAAAPAVPVRCFTFC
22000	33174	^	22931	ľ	400	FTDIVIMPKRKSPENTEGKDGS
						KVTKOEPTRRSARLSAKPAPPK
		l		1		PEPKPRKTSAKKEPGAKISRGA
						KGKKEEKQEAGKEGTAPSENG
ļ		1				
	ŀ				1	ET\KAEEIHISRSTVNVSTSRGTP
		١.	22222		205	PSTLSVKGQIETVRVKGTEN
22807	53175	A	22938	3	285	SSCPAREQLQRQRQHPAAAPAV
					ŀ	PVRCFTFCFTDIVIMPKRKSPEN
				l		TEGKDGSKVTKQEPTRRSARLS
1		l	1	l		/AVFV*FKLSLQKPAPPKPEPKP
		_				RKTSA
22808	53176	Α	22939	1	413	
22809	53177	Α	22940	2	454	AEVQVEEAGLRRALRVSSCSGR
		l				GSIQRRFYAVPVRCFTFCFTDIV
						IMPKRKSPENTEGIDGS\KVTKQ
						EPTRRSARLSAKPAPPNPEPQPR
						KTSAKKEPG\AKISRGAKGKKE
ŀ	İ					EKQEAGK\EGTAPAENDETKAE
						EAQKTESVDNEGRMNCP
22810	53178	В	22941	55	186	
22811	53179	A	22942	2	397	
22812	53180	В	22943	339	470	
22813	53181	Α	22944	1	139	
22814	53182	Α	22945	1	416	MPLLYHLSRCAYVQQCNFANP
				1	l	VVSALCTCLVKPVVEKCTPYA
l	1	1	1		I	VLLEALALRNVRQLEDLVIEAV
	1	1	i		I	YADVLRGSMDQRNQRLEVDYS
	1	1	l	1	I	IGRDIQRQDLSAIARTLQEWG/C
	Ì				1	HQHLLSLRSAASQRQQCGYHK
		1				LHDKQRLLRG
22815	53183	Α	22946	315	570	

SEQ ID NO:	of peptide		SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22816	53184	Α	22947	933	1841	RPAEPDPGSGHPEPKSPTLSAV
						MSAEVKVTGRNQEQFLLLAKS
	1					A\KGAVLATLIHQVLE\APGVY
						VFGELLDMPNVRELAESDFAST
						FRLLTVFAYGTYADYLAEARN
						LPPLTEA\QKNKLSDTFSVV\TL
						AAKVKCIPYAVLLEALAL\R\NV
						RQLEDLVIEAVYADVLRGSLDQ
						RNKRL\EV\DYSIGRDIQRQDLS
	1					AIARTLQEWCVGCEVVLSGIEE
	İ					QVSRANQHKEQQL\GLKHQIES
						EVANLKKTIKVTTAAA\AAATS
			١,			QDPEQHLTELREPAPG\TNQRQ
						P\SKKASKGKGLRGSAKIW\SKS
22817	53185	A.	22948	I	1319	
22818	53186	Α	22949	365	1331	RAGGGAPWRFTGTFATVAPET
	1				1	AADLSLHRHLLSRTTSAWGTA
						CITVPIEPPTWPTWAVPIQWGG
					1	SATCIQEDKLRQGILYGYSQQV
						NQAMDSGLRP\VVDSVNAIRVP
				1		QDYVTQSVPLREMNGSLGVLA
						QQLQNAKLQADAAHSALKQSD
	1					DLKPVFDQAFTKVVTTPADAL
						QPLIPAAQTFTQQLVMVGDYIA
	1				ŀ	QQGTQECGKTFSRASYLVQHS
	İ					RIHIGKKPYECKECGKAFSSGS
	1					YLVQHQRIHTGEKPYECNKCG
						KAFTVYGQLIGHQSVHTGEKPF
				i		ECKECGKAFRLNSFLTEHQRVH
						TGEKPFKCKKCGKTFRYSSAP
22819	53187	Α	22950	2	206	GGGQGATSFGRPESGFIRHQPPP
	l					PCRRSSTPTRSKLYTL\SAPE*RS
						VPTSAPGPQDRPPGSVSKKSW
22820	53188	Α	22951	571	1102	MGSVSLGNPQGPFQFTRLPLAN
						VIHPTFFWETGPRGADSLGAKG
		l				RKWQPTFTSRCNLKVYLGRAG
						SQAATGGPSPSLPNLHVKKLRP
						RKAEAWPGRRNTGKVSGR/LA
		l				ALRAPYGESFARAAGPRDAPS/
						QQPRPQPEHAPTLISLGSNFGG
						MVEESWCRMNPDSGRPKEVAP
						WPPPSTK
2282 I	53189	A	22952	1	591	MGPHWKDFKEIDVKKYIDALSI
	1	İ		1		HTYTHTHTHADITDGDHGG
	1		1	I		GALLELEINSHLKVQFRELNITA
		l	1	l		AKETEDGGGGKAIIIFVPFPQLK
				l		SFQKIQVWLRRILPKPTRKSCT\
				1		QNMHWPPRSRTLKAMRDSIPE
						DLVF\PSEIMCMRIHVKLDGSRL
				-		IKAHLDKAQQYHVERNAETFS .
	1	İ				GVYMKLTGKDINVEFPEFQL

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hed	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22822	53190	Λ	22953	2	508	APPFELPRAAGIRHEGKFSQEK AMFSSSAKIVKPNGEKPDEFES GISQALLELEMNSDLKAQLREL NITAAKEIEVGGGRKAIIIFVPVP QLKSFQKIQVRLVRELEKKFSG KHVVFIAHRRILPKFTRKSRTK NKQKKPRSRTLTAVHDAIPEDL VFPSEIVGKRNR
22823	53191	Α	22954	145	395	
22824	53192	Α	22955	536	717	
22825	53193	A	22956	1	729	
22826	53194	A	22957	1	1158	
22827	53195	A	22958		2869	MAHLRSCCSSLYQPGHLLPPHN PQDHPLTAPSLTHLTRRAYFTD EKTKGGQSAYAVSLLRECVKL RYSDPTVPLMAAK VCIGSLRW MAAKHLAGVLIHSLSECYWS PLSHPLPEFMGKEESSFATQAL RKPHLYEGDNLYCPKDNIEEAL LLLIESSMLANKLLLLEMEQ QSPGVESGVDRQLSPYSDTNVN AQNAQIVPVAAKSAQNEVPGY PEETEKSRYQSNLITGPGDLRL LLAAPECLPPRMSPPS
22828	53196	В	22959	28	139	
22829	53197	A	22960	60	812	LASPFVPLHRCSAGSOPPCPVP EGLIRIYSMRFCPYSHRTRLVLK AKDIRHEVVNINLRNKPEWYY TKHPFGHIPVLKTSKCQLIYES VIACEYLDDA YPIGRKLFPYDP YERVRQKMILELFGKVPHLPR EFPGSVEDVGENAPNLKAALR\ QEFSNLEEILEYQNTTFFGGTCI SMIDYLETLWFWFERLLECMGY WNCVSHTPALRLWISAMKWD PTVC/ALLMDKEHF\QGFLNPYF QNNPNAFDFGLC
22830	53198	A	22961	2	251	ILLDFKPWDDETDMAQLEACV CSIQLEELV*GASKLVPMGYGI RKLQIQCVVEDDKVGTDLLEEI TKFEEHVQSVDIAAFNKI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		İ		sequence		
	Ļ	<u> </u>				
22831	53199	Α	22962	353	1071	APLWPRAERKFYKQMNGPVAS
				1		ASRQSSAPG/NDHSELVLQIASL
1		l				EVENQSLRGVVQELQQAVSNL
		l		1		EAWLN\IRREKSSPGH/RSTVP
		l				QTQHVSPMRQVERPSPRPPYYD
		l		l		PGQEAEDNEDDDIDLDNKEED
1		1		ľ		KEARRLREERLRLYAKKAKNP
l				1		SLVAKSSILLDIKPWDNKTDMT
	1.0	l		l		QLEACVRSIQLDGLVSGASKLA
		l				SVGYGIRKMQIQCVVEDDKVG
l						TDLLEEEITKFEEHVQSVDIAAL
22832	53200	Α	22963	3	482	PWANEAPAKK/PATPAEG/DAR
		l		l		DDDIDLF\GSDNEE\EDKQAGHS
1				l		CGEER\LR\HYAGEEGPRKPAL/
		1		l		VWPKSLPSLLDVK\PWDDDDGP
		1		l		WPSLEACVR\SIQLDGLVWGGF
1		l	l	l		PSLVPVG\YGIRK\LQ\IQCVV\ED
		l		l		DKVGTDFPWRREIT\KFEEARA
		1				RSVDIASFSTRS
22833	53201	Α	22964	1	2835	
22834	53202	В	22965	571	1337	
22835	53203	Α	22966	Į.	489	
22836	53204	Α	22967	1	618	
22837	53205	Λ	22968	3	5227	SWSRVRPRTRPEGLRQLKMRT
		1				GWATPRRPAGLLMLLFWFFDL
				l		AEPSGRAVN\DPLTIFHGNTGKC
		1		l		IKPGYGWIVADHCDETEDKLW
		1		l		KWVPQHRLFHLHSQKCLGLDIT
		1				KSVNELRMFSCDSSAMLWWKC
1				ł		EHHSLYGAARYRLALKDGHGT
		ı		l		AISNASDVWKKGGSEESLCDOP
		l		l		YHEIYTRDGNSYGRPCEFPFLID
		1		l		GTWHHDCILDEDHNGPWCATT
		1		1		LNYEYDRKWGICLKPENGCED
						NWEKNEOFGRCYOFNTOT
22838	53206	A	22969	3	453	GOSPILVIYHETKRPSGISERFSG
		1		ľ		STSGNTATLTITGTOAMDEADY
		1		1		YCQAGRDFGGGTKLTVLGQPK
		l	l	[		AAPSVTLFPPSSEELQANKATL
	1		l	1		VCLISDFYPGAVTVAWKAHSN
1		l	ŀ	[		PVKAGVETTTPSKQSNNKYAA
		1	ĺ	l		SSYLSLTPEO*KSHRSYSC
22839	53207	A	22970	2	354	30 LEST LEG. VOLVO 19C
22840	53207	IA.	22970	1	315	
22840	22208	IA.	229/1	1 .	313	

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X-Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
22841	53209	A	22972	I	712	MAWTLIFITLLTOGTGSWAGA LTOPASVSGSPOGSITISCIGS SDIGAYYFVSWYQQYPGKAPK LLIFDVSGRPSGISSRFSGSKSGN TASLTISGLQAEDEADYYCASY ARSSTVVFGGGTSVTVLGQPKA APSVTLYPPSSEELQANKATLV CLISDFYPGAVVAWKADGSP VNAGVETTTPSKQSNNKYAAS SYYLSLTPEQWKSHRSYSCQV THEGSTVEKTVAPTECS
22842	53210	Α	22973	1	1779	
22843	53211	Α	22975	I	2001	
22844	53212	A	22976	292	1497	TGQLNGFHEAFIEGTTEITSES VGGEHPIDKICDQISDAVLDA HLQQDPDAKVAICETVAKTGMI LLIAGEITSRAAVULPRKVVRE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE LEDENTROLKEENIKAVVPAKY LDEDITYHLQPSG/RTFVIGGPQ GDAGLTGGRKINDTYGGGQA HGGIGAFSGKDYTKVDRSAAY AARWYAKSLVKGGLCRRVLV OXSYAIGVSHPLSISIFHYGTSQ KSERELL*IVKKNFDLRPGYVR DLDLKKRIYGRAAYGHFGRDS FPWEYPKKLIY

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
W				sequence		
22845	53213	I <sub>A</sub>	22977	67	1579	FFVRSLSHTDTNMNGQLNGFH
22843	33213	A	22911	07	1379	EALIEEGTFLFTSKADGEGHPD
	,					
						KICDQISEAVLDAHLHQDPDAK
					i	VACETVAKTGMILLAGEITSRA
						AVDYQKVVREAVKHIGYDDSS
						KGFD\YKTCNVLVALEQQSPDI
						AQGVHLDRNEEDIGAGDQGLM
		l			l	FGYATDETEECMPLTIVLAHKL
		1	0			NAKLAELRRNGTLPWLRPDSK
		1			l	TQVTVQYMQDRGAVLPIRVHTI
		ł				VISVQHDEEVCLDEMRDALKE
						KVIKAVVPAKYLDEDTIYHLQP
		1				SG\RFVIGGPQGDAGLTGRIIIVD
						PYDGWGAHEEGAFSGKDYTKV
		1				\SRSAAYGARWVARSLVKGGL
	1	1			ŀ	CRRVLVQVPGPPKKPDEDLYLE
	1				ļ	CEPDPVLALTQTLSFQVLMPSG
					l .	PLPRTSVVPREQQMLPLKDSDL
	1				1	LTQPWYSGNCDRYAVESALLH
						LQKDGAYTVRPSSGPHGSQPFT
						LAVLLRGRVFNIPIRRLDGGRH
		1				YALGREGRNREELFSSVAAMV
						QHFMWHPLPLVDRHSGSRELT
22846	53214	Α	22978	3	325	
22847 22848	53215	A	22979	7	1014 592	LI COODUCE OCTAVOR OF THE
22848	33216	l^	22980	ľ	1392	LLCSGRHSLQGVMGILSFLPVL ATESDWADCKSPQPWGHMLL
					i	WTAVLSLAPVAGTPAAPPKAV
		1				
		1				LKLEPQWINVLQEDSVTLTCRG THSPESDSIQWFHNGNLIPTHTQ
	ŀ	1	i			
		1				PSYRFKANNNDSGEYTCQTGQ
		1				TSLSDPVHLTVLSGQWRKAPG
		1				WTWEGQDG*NLLSGRGLQERG
22040	53217	١.	22981	32	1150	VACLLGSIAVSCLSTYQWLFLP
22849	33217	Α	22981	32	1150	EKAVTAVLWAPARLQGVMGIL
		1				SFLPVLATESDWADCKSPQPW GHM\LLW\TAVLFLAPVAGTPA
		l				APPKAVLKL\EPQW\INGGSRED
		1				
						SVTL\TC\RGTHSP\ESDSIQWFH
						NG\NL\IPTHTQPSLQVSRANNN\
		1				DSGE\YTCQTGQTSLSDPVHL\T
		1				VLSEWLVLQTPHLEFQEGET\IV
	1				ĺ	LRCHS\WRDKP\LVKVTFF\HNG
		1				KSKKFSRSDPNFSIPQANHSHSG
		1				DYH\CTG\NIG\YTLLFIPRLLTIT
						VQAPSSSPMGIIVAVVTGTAVA
	1	1				AIVAAVVALIYCRKKRISANPT
				1	Ì	NPDEADKVG\AE\NTITYFTSQW
	1					HPGWLWEEPDDQNRILVSIVLA
1	1	1				LGIWRRNSEREDLVFPGLKFPL
						GEDTGRCCQFPKEKVFLPGVIL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	cudon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
l				sequence		
22850	53218	A	22982	1	471	MKSSFOGHNKEKREKKGPSLFS
22000	55210	ľ	2	ľ		AAYGGGSCLLLSIMAALRPLVK
		l			i	PKIVKKRTKEFIWHQSDQHVKI
l						KCNWRKPRG\QILMPNTGHGSN
į.		l	1			/KKTKHMLPSGFRKFLVPSVKE
Į.		1				LEVLMCNKSPCAEIAHSVSSK\N
			l			KTIVERAAQ/LAVRVTNPNARL
			i			RSEENE
20051	50010	١	22082	102	564	
22851	53219	A	22983	103	364	GIFRGIMAALRPLVKPKIVKKR
						TKKFIRHQSDRYVKIKRNWRKP
			l			RGIDNRVRRRFKGQI\LMPNIGY
1						G\SNKKTK\HMLPKWLSGKFLG
		l			ľ	SHQRSREL/EKLLA*CCNKFFTC
l				1		CPRFVSHCFSFQGTGKGLRGER
		<u>l</u>				SLPNWAFRVTNPNARLRSEENE
22852	53220	Α	22984	3	177	
22853	53221	Α	22985	74	735	RFAGAGAIPEARARPPDVQAAE
		ł				EEKEMDLPDSASRVF/CGDASS
		į.				NMGEHRMNVNAIIL\AQKNML
		1				D\RFEKT\NEMLLNFNNLSSARL
			İ			ASRLSERFLHHTRTLVEMKRDL
		l	l			DSIFRRIRTLKGKLAROHPEAF
						SHIPEASFLEEED\EDPIPTOAPT
		į.	ŀ			TTIATSE\QSTG\SC\DTSPDTV\S
	1	i i	ŀ			PSLEPLASEDLSHVQALASPAI\
ŀ		t		İ		NGPOPDOIDEEMTGRIALPARC
22854	53222	A	22986	3	275	MRTVAEKVDAVRIVHIHSVI\IL
22034	33222	n	22,00	ľ	273	RRSDKRKDRVEISPEOLSAAST
		l	ł			EAERLAELTGRPMRVVGWYHS
l		1				HPHITVWPSHVGKYHGVAMFI
i		1				VFRN
22855	53223	A	22987	1	886	AKMAVOVVOAVOAVHLESDA
22833	33223	^	22901	l <sup>*</sup>	000	FLVCLNHALSTEKEEVMGLCIG
1				l		
						ELNDDTRSDSKFAYTGTEMRT
						VAEKVDAVRIVHIHSVIILRRSD
1				i		KRKDRVEISPEQLSAASTEAER
l						LAELTGRPMRVVGWYHSHPHI
						TVWPSHVDVRTQAMYQMMDQ
		l		1		GFVGLIFSCFIEDKNTKTGRVLY
		1		1		TCFQSIQAQKSSEYERIEIP\IHIV
1		1	l	1		PHVTIGKVCLESAVELPKILCP/
		l	l	1	l	QEEQDAYRRIHSLTHLDSVTKI
				1		HNGSVFTKNLCSQMSAVSGPLL
				1		QWLEDRLEQNQQHLQEL\HQE
				1		KEELMQELSSLE
						· · · · · · · · · · · · · · · · · · ·

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22856	53224	A	22988	110	988	LKAMSWAKMAVQVVQAVQA
22050	John I	Ι.	22700	1.*		VHLESDAFLVCLNHALSTEKER
				l		VMGLCIGELNDDTSRSDSKFAY
				1		TGTEMRTVAEKVDAVRIVHIHS
l				i		VIIL\RRSDKRKDQVEISPGQLSA
				ı		ASTEAEKLAELTGHPMRVVGW
						YHSHPHITVWPSHVDVRTQAM
				ı		YOMMDOGFVGLIFSCFIEDKNT
				1		RAGRVLYTCFQSIQAQKSSESL
l				1		HGPODFWSSSQHISIEGQKEEER
		1		1		YERIEIPIHIVPHVTIGKACLESA
				1		VELPKILCQEEQDAYWRIHSLT
		1		i		HLDSVTKIHNGSVFTKNLCSQM
1		1				SAVSGPLLQWLEDRLEQNQQH
						LQELQQEKEELMQELSSLE
22857	53225	Α	22989	1	1362	
22858	53226	A	22990	284	542	KLFQKKNKKSSCISDNFFFLLAI
	!	1		I		MVNLLQIVRDHW\VH\VLVPM
		1	i			GFVIGCYLDRKSDERLTAFRNK
		4_				SMLFKSFPHRELQPSEEVTWK
22859	53227	Α	22991	272	1241	FPGGAACDICWVKEENGSLMS
		1				MKMGWKAKSCLPVTVHPRCL
		1				TPYSARLSSTFPL*NSITTGP*QS
				1		PACKRPF*LTTC*GGSRRNSNR
				1		KAA*GPCSPPPPSPPPSPAT\LPR
l		1				GPAGLQPPGVPVLPPLRPRKHY
		1				APGGLP\PPASLLEDDDDTFCTS
		1		1		QAMQPTAPTKLSPPALLPEKDS FSSALDEIEELCPTSTSTEAATA
		1		1		ATDSVKGTSSEAGTOKLDGPOE
		1		1		SRADDSKLMDSLPGNFEITTST
ì		1		1		GFLTDLTLDDILFADIDTSMYDE
				1		DPCTSSSGTASKMAPVSADDLL
				1		KTLAPYSSQPVTPSQPFKMDLT
			ł	1		ELDHIMEVLVGS
22860	53228	A	22992	1	3234	MELKCVCLKNRSPSAEEQPSLL
		1		1		PEPSAQESHGGHPRSALYGRER
i	1	1		1		LTGOPPSSPHPNPAPWASCPAVI
			ł	1		DPVPIYSSVAEAYDSGTEGIPTV
			ł	1		PGLR\HVESGSLRATAQRLPPER
			ł	1		RV/OORTTAPPPROPPRRPLRAS
l		1				LSAAPPEPREPVVAAGPEPGSL
1	1	1	l			QRPARTALPNRPEPQAASPGTL
1	1	1	1	1		VEKYAAPVATQDLLDENPCTS
1		1	1	1		RLGVVDLQAVLTLRKLENLRS
			l	i		GAQTPLEGGQSGRVGPHALRW
		1		1		NRGEEEGERLLC
22861	53229	A	22993	307	1906	
22862	53230	Α	22994	1	1752	
22863	53231	Α	22995	588	704	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nuclcotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22864	53232	A	22996	3219	3407	DQVVWTKRLQDAILVLV*EFRL HSPALRLCVMKQVGMSQGELE
22265	50000	١.	22007		588	WGQFLKLSSRNRKRSGERI
22865	53233	A	22997	1	917	
22867	53235	A	22999	1	1692	
22868	53236	A	23000	1	2266	MKRIRDYYERSHANNLDNLEE
22000	33230	^	23000	l'	2200	MDELLETHNLPGLNHEETENLN
1		1				KPITTSLSPLAGVSSTKKNTKNE
						LSIVLPFTGSILPVGFATRPRGL
i	ŀ			l	i	ASSNPILLHMVAFLFANGTEEL
			i		PVPLMTPTRQKKVIPLIPLMVG	
	1				LGVSASTIALGTGITGISTSVTTF	
i						RSLSNDFSASITDISQTLSVLQA
ŀ						LVDSSAAVVLQGQLRGLDLLT
	ł					AEKGGLCIFLNEECCFYLNQSG
	ĺ	1		i		LVYDNIKKLKDRAQKLANQAR
		1		l		QNLDSMLHGTGMKSDSDQKKS
l		1		i		ENGVTLAPEDTLPFLKCYCSGH
		1				CPDDAINNTCITNGHCFAIIEED
	l .	1				DQGETTLASGCMKYEGSDFQC
l	1					KDSPKAHVRRTIECCRTNLCNQ
		1			ļ	YLQPTLPPVVIGPFFDGSIRWLV
						LLISMAVCIIA\MIISSSCFCYKH
	l				1	YCKSISSRRRYNRDLEQDEAFIP
						VGESLKDLIDQSQSSGSGSGLPL
		1				LVQRTIAKQIQMVRQVGKGRY
		1				GEVWMGKWRGEKVAVKVFFT
						TEEASWFRETEIYQTVLMRHEN
		1				ILGFIAADIKGTGSWTQLYLITD
ļ						YHENGSLYDFLKCATLDTRALL
	į	ŀ			l	KLAYSAACGLCHLHTEIYGTQG
1	1					KPAIAHRDLKSKNILIKKNGSCC
į.						IADLGLAVKFN\SDTNEVDVP\L
						NTRVGTK\RYMAPEVLDESLNK
1		1		l		NHFQPYIMADIYSFGLIIWEMA
						RRCITGGIVEEYQLPYYNMVPS
						DPSYEDMREVVCVKRLRPIVSN
22869	53237	Α	23001	68	636	GVTSNPKALLGALIAPAFASLPE
						GRGKRIQALNMFRPKNSWRLG
I			1	1	1	LKYPPSSKEKTQVPKTLISGLPE
1	1	1	1	I	I	RKSTTRVGEKLQSAHKMPLSPG
						LLLLLFSGATATA/APLPLEGGP
1	1	1	I	I		TG\RDSEHMQEAAGIRKSSLLTF
I	I	1		1	[	LAWWFEWTS\QASAGPLIGKEA
				1		REVARRQEGAPP\QQSARRDRM
						PCRNFFWKTFSSCK
22870	53238	A	23002	1	144	GIRLPSSADWQKTEAQKRREQL
1	1	1		I	1	ILLDELVALVNKRDALVRDLDA
	1	L		I	l	QEKQ

W 0					T	
SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	noa	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	sequence		02/340,217	sequence	or pepade sequence	l l l l l l l l l l l l l l l l l l l
22871	53239	Α	23003	2	667	RWNSSDPDADRTTLNHADHSS
			1			KIVQHRLLSRQEELKERARVLL
			1			EOARRDAALKAGNKHNTNTAT
						PFCNROLSDOODEERRROLRER
				1		AROLIAEARSGVKMSELPP\YG
						EMAAEKLKERSKASGEONSKL
				ł		VDLKLKKLLEVQPQVANSPSSA
	1					AQKAVTESSEQDMKSGTEDLR
1		ł				
ŀ		ŀ		1		TERLQKTTERFRNPVVFSKDST
		l				VRKTQLQSFSQYIENRPEMKRQ
		-			<u> </u>	RSIQED
22872	53240	Α	23004	1	1437	
22873	53241	Α	23005	1	2009	MAKPYLYKKYGNQQGVAAIPD
						VPAAWEAEPSPIPSPVLGRKPN
l	1		1			ASQSLLVWCKEVTKNYRGVKI
	i					TNFTTSWRNGLSFCAILHHFRP
	l		1			DLIDYKSLNPQDIKENNKKAYD
	1					GFASIGISRLLEPSDMVLLAIPD
į		1				KLTVMTYLYQIRAHFSGQELN
ŀ	1	1				VVQIEENSSKSTYKVGNYETDT
		1			l	NSSVDOEKFYAELSDLKREPEL
1		l			1	QQPISGAVDFLSQDDSVFVNDS
		1				GVGESESEHOTPDDHLSPSTAS
		1				PYCRRTKSDTEPQKSQQSSGRT
		ĺ				SGSDDPGICSNTDSTQAQVLLG
		ı				KKRLLKAETLELSDLYVSDKK
1		ł				KDMSPPFICEETDEOKLOTLDIG
						SNLEKEKLENSRSLECRSDPESP
l			1			IKKTSLSPTSKLGYSYSRDLDLA
						KKKHASLRQTESDPDADRTTL
			1			
						NHADH\SSKIVQQDEERRRQLR
					1	ERARQLIAEARSGVKMSELPSY
	1		1		1	GEMAAEKLKERSKASGDENDN
			1		l .	IEIDTNEEIPEGFVVGGGDELTN
	İ		1			LENDLDTPEQNSKLVDLKLKKL
		I	1		1	LEVQPQVANSPSSAAQKAVTES
	1	1	1		1	SEQDMKSGTEDLRTERLQKTTE
		I	1	l	1	RFRNPVVFSKDSTVRKTQLQSF
		I				SQYIENRPEMKRQRSIQEDTKK
l		I				GNEEKAAITETQRKPSEDEKGF
1	1	I		l	1	KDTSQYVVGELAALENEQKQI
1	1	I	1	i	1	DTRAALVEKRLRYLMDTGQNL
ŀ	1			1		DITUTAL PERIOD OF ALL

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	liou	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22874	53242	A	23006	145	3217	LDEYEARLTLANLDDFEEDNEI
						DDENRVNQEEKAAKITELINKI
						NFLDEAEKDLATVNSNPFDDPI
			ŀ			AAELNPFGDPDSEEPITETASPR
			1	1		KTEDSFYNNSYNPFKE\VQTPQ
						YLNPFDEPEAFVTIKDSPPQSTK
				1		RKNIRPVDMSKYLYADSSKTEI
						EELDESNPFYEPKSTPPPNNLVI
						PVQELETERRVKRKAPAPPVLS
						PKTGVLNENTVSAGKDLSTSPK
				1		PSPIPSPVLGRKPNASQSLLVW
						KEVTKNYR
22875	53243	Α	23007	1	165	RPAAPRHERRPAADHGHHRPE
				]		GRGTGQKACSELPSDEACS/GIA
		_				IG*SLLCSACSVRSRKRQW
22876	53244	В	23008	26	625	
22877	53245	A	23009	325	526	LI CUMPLICITIVINI ICENTA ICENT
22878	53246	A	23010	1946	2224	LLQWRHGLLNPINSPYNSPILPV
						QKPDKPYRLVQHLCLIYQIVLP
						HPMVPNPYTLLSSIPFTTHYSVI
		l				DLKHAFFTIPLYPSSQPLFAFTW
		l				TDPDTHQAQQITLAVLPQCFID
		l				PHYFSQAQISSSSVTYLGIILMK THIGLGAVEOGVVLVGEARAA
			ĺ			QEPMEWVGGSGMVGCRSRAL
		l		[	ŀ	RGKAAKARREIERSAVTIVPVL
		l				DENPARHIIPDTTPDHHDCISLIF
		1	l			LTFTPFPHISFFPVPHLEHTWFIE
						GSSTR
22879	53247	A	23011	1	1479	MEENFIRVFFTTTTMLLLIALIK
22017	33247	^	23011	ľ	14/7	OGELCESFDGKSLDGRDERVVI
		1	1			CFDAAAGLHRACSIRGEOONL
		l			1	GTHYFTWQRLFAEVIKLGGLM
	1	1				KPKARNASSLWKEKTKNGFFS
						LETPEERQTSDLHKYQLAFRLF
		l				FIKYKNPAQFMAHLATTLRHF1
		l				ALDPESSHNFONLFSSSHLMHII
						SAPRLLRLYSRFVESPTVTIVPG
						PDFNPASNIIPDTTPDPHDCISLI
			1			HLAFTPFPYISFFPVPHPDHTWF
						IDGSSTRPNCHTPAKAGYAIVS
						TSIIEATALPPSTTSRQAELIALT
						QALTLAKGLRINIYTD/SHILHH
						HAVLWA/E/RGFLTMQGSSIINA
						SLIKTLLQAALLPKEAEVIHCKO
		1				HOKASDPITOGNAYADKVPSL
		1		l		AHAPTLTLAPVLQCPLREVAGT
		1		1		EDDNGKAVLEIPSNIRKSAHVP
		1	1			VOTWKGWDPTNLLGGWFSNL
		l	1			GGFKTLVGTVIFIIGFLLFLPCVI
				1		

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22881	53249	A	23013	27	496	
22882	53250	A	23014	117	210	
22883	53251	A	23015	124	414	
22884	53252	A	23016	3	627	
22885	53253	Α	23017	1	567	
22886	53254	Α	23018	44	551	GGGRSAPLQLGFMSRQAKDD FLRHYTVSDPRTHPKGYTEYKV TAQFISKKDPEDVKEVVVWKR YSDYRKYLHGD/LAYTHRYLFIR RLEEFPAFRAQ\VFGRIVEASV DSRERYRKGAEDLLRFTVHIPAL NNSPQLKEFFRGGEVTRPILEVS RDLHILPPPLIPTPPPG
22887	53255	A	23019	25	488	
22888	53256	c	23020	9	119	
22889	53257	Ā	23021	52	241	
22890	53258	В	23022	4	402	
22891	53259	A	23023	249	984	GSTMYKMNICNKPSNKTAPEK SVWTAPAQPSGPSPELQGQRSR RNGWSWPPPHPLQIVAWLLYLF FAVIGFGILVPLLPHHWVPAQY AVSLGKRRAGPWNPFLSVSAFT ILPQCMGAIFAGHLVVHLTAVS IDPADANVRDKSYAGPLPIFNR SQHAHVIEDILCNLCNVDVSA RSKHCSACNKCVCGFDHHCKW LNNCVGERNYRLFLHSVASALL LNNCVGERNYRLFLHSVASALL CYLLLV/LGGHICLRGVLCOPH ASAHQPTL
22893	53261	A	23025	[11]	1368	OODSGSG ALADGSGI DDDDDGDI
22893	53261	A	23025		1368	QOPSGGALARGSGLPPRFGRL PARSPVOKRKGGWPKGKKKGGWKGKKKGWGWKGKWF PRDLAANRVPTTGYMIELNEQRS QLRATHPDLPFTEIMKILAVQ WAQLSQDKKGRKYAGKALGV TPPPSALTPVALIEEPKQLDIQSL GMHTVTEGLAGRQVILQQVQ GGAWESALTFVALIEEPKQLDIQSL GGAWESAFLTCPWAMAAAAP GPTLTMLQPSISAVGFSQMAMG PHVQGRASKLSIYVYPRRQLST TTAMGDELNDLYCGTSRGUFSS LCNKKEPAAKAAPAAPLSSPGP PALGHAGHGRGULLERSLTLFH LYELLKYPRASPGLWPPRVWM PHSLTCGTFWVPMIMMRFCAH VASGLLRPPROVYAHLSPAQG RHPGENGIIVTSRAAVRIRCSS CSRSLTGNLKKILPRTQSTWSL WRKRROK

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22894	53262	Α	23026	2	405	
22895	53263	Α	23027	1	391	
22896	53264	Α	23028	3	463	
22897	53265	Α	23029	74	466	FAFNMPEPAKSAPAPKKGSKK
		l				AVTKAQKKDGKKRKRSRKESY
		l				SVYVYKVLKQVHPDTGISSKA
		l			0.0	MGIMNSFVNDIFERIAGEASRL
	İ					AHYNKRSTITSREIQTAVRLLLP
		_				G\EL\AKHAVSEGTK\AVTKYHQ
22898	53266	Α	23030	168	597	KEETDFLFWFCHYCFIMPELAK
			ŀ			SAPAPKKGSKKAVTKAQKKDG
						KKRKRSRKESYSVYVYKVLKQ
1		1				VHVPDTGISSKAMGIMNSFVTTF
		l	1	ļ		FE\RIAGEASRLAHYNKRSTITS
1		l				REIQTAVRLLLPGELAKHA\VSE
		_				GTNA\VTKYTSSK
22899	53267	Α	23031	18	321	FTATWLNQEVGIVGKYRTRYG
			l			VSLQKMVKKIEISQYAKYICSFL
						WQNSKMKRRAVGIWCCG/SCM
				l		KTVKVSSSRNHIVWTYNTTSAV
	*****	Ļ.				TVKSTIRRLKQLKDQ
22900	53268	A	23032	12	445	FTCRSHVSEVEERCVEAPGLLG
						SRSITGFSLHSKQAKRTKKVGIV
	1	1	ļ			\GKYGNPLWGPSLR\KMVKK\IE
ł		1	1			ISQHAQVHFALFCGKTKMKRR
			l			AVGIWHCGSCMKTV\AGGAWT
	ĺ		i			\YNTT\SAVTVKSAIRRLKEFERP
22901	53269	١	23033	3	175	SRTLLYSLRTSLGL
22901	53270	A C	23033	378	434	
22902	53270	A	23034	1	578	MVRCVRLVEAGSVVRYLSTSIC
22903	332/1	A	23033	1	3/8	
					1	RPVVDAGSRALCLQEWADSQQ VKEKQYSSRDVQRAAALNIYRI
						PPSSRKPALCPTPRDRLEYDEDR
						LEHIAYVRARELHTLEVTGLET
					ŀ	VAOSKAHVASLEGLIPEDKVVL
	1					LAGSPLQNEATLGQCGVEALTT
						LEVVGRRLGASL/HTSASKHTM
22904	53272	С	23037	236	379	VRALTYCSSREETFTAI
22904	53272	A	23037	178	534	VVQKTEPRLCFAVSPPFSCSLPL
22905	33213	^	23038	1/0	334	PSLFPHSPLLGRPKRQAKPAAD
İ						EGFWDCSVCTFRNSAEAFKCSI
l						CDVRKGTSTRYLKICVTWVVV
			1			VI*QVFIEHTAICALGVQWYKD
L	L	_				RPGPCSYGA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
22906	53274	Α	23039	44	1012	VGGGGVQPSMTMGDKKSPTRP
i						KRQAKPAADEGFWDCSVCTFR
	l			ŀ		NSAEAFKCSICD\VRKGTSTRKP
	1					RINSQLVAQQVAQQYATPPPPK
1						KEKKEKVEKAGQKRKPEKRPR
	1					KFSP/SVVTKKNTNKKTKPKSD\
		l				ILKDPPS\EANSIQSANATTKTSE
	1					TNHT\SRPRLEKRGTGGIAQQFG
						QLTVG\NVTVIITDFK\EKTRSSS\
						TSSSTVTS\SAGS\EQQNQRQLR
						GFRGATDKGSLPVPPNPKGRPC
						QAVNDESFLKLAHGNCGKLWN
1						SGVWKFQNLPPAHACLHPWEN
1						LSCGTSTSLVDCCQDNFLLCHG
						HSWPPRDFRHPDDYSGHF
22907	53275	À	23040	1	626	MAGCRNPCKAESHRSKSLLSEA
	l					ESLFREAESQHDRGSKNAGHR
	l					DESVEVWLSLLSSSQTRAEIER
1	l					QSRGKQWLRHLLRDTEGPKET
						VRPELATRAAMDVAEVEFLAE
	1					KELVTIIPNFSLDKSYLIGVDNIP
	İ			i		KADEIRTLVKDIWDTRIAKLRV
	Ì					SADSFVRQQEAHAKPDNLTLM
				Į.		EINPSGTFLTQALNHTYK/LRTN
						LQPSESTQSQDF
22908	53276	Α	23041	1	951	MDAAEVEFLAEKELVTIIPNFSL
						DKIYLIGTWELKYREIRWLSQG
						PHGKPHADPGSSILQNSSGPSG
						VRTNGKSKPWEPFVAEEFAHQ
	l l					FHESVLQSTQKALQKHKGSVA
						VLSAEQNHKVDTSVHYNIPELQ
	1				:	SSSRAPPPQHNGQQEPPTARKG
	1		İ			PPTQELDRDSEQEQEEDDEDGE
	ŀ					DEEEVPKRKWQGIEAVFEAYQ
						EHIEERTLERNRSSTAADKAQL
				i		LAANRQGHGQPTDPASDNIPKA
	1			Į.		DEIRTLVKDMWDTRIAKLRVS
				l		ADSF\VRQQEAHAKL\DNLT\LM
						EIHTSG\TFLTQALNHMYKLRT
						NLQPLESTQSQDF
22909	53277	Α	23042	1	626	MAGCRNPCKAESHRSKSLLSEA
						ESLFREAESQHDRGSKNAGHR
	1		l			DESVEVWLSLLSSSQTRAEIER
	1					QSRGKQWLRHLLRDTEGPKET
	1		İ			VRPELATRAAMDVAEVEFLAE
	1		1			KELVTIIPNFSLDKSYLIGVDNIP
	1		l			KADEIRTLVKDIWDTRIAKLRV
	1		l			SADSFVRQQEAHAKPDNLTLM
	1		l	l		EINPSGTFLTQALNHTYK/LRTN
1	1		l		1	LQPSESTQSQDF

SEO ID	SEO ID NO:	Ta 1 - 1	SEO ID NO:	N	North did booting of last	Amino acid sequence ( X=Unknown.
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1.0.	sequence		09/540.217	codon for pentide	of peptide sequence	deletion, \=possible nucleotide insertion)
	,			sequence		·
		<u> </u>				
22910	53278	Α	23043	2	642	RPRRLLRDAEGPEETVRLWPAA
					İ	RAAMDAAEVEFLAEKELVTIIP
i	1					NFSLDKIYLIGGDLGPFNPGLPV
					1	EVPLWLAINLKQRQKCRLLPPE
l						WMDVEKLEKMRDHE\RKEETF
		1				TPMPKPFTTLELT\KLLLNHASD
						NIPKQDEIRTLVKDMWDTRIAK
						LRVSADSF\VRQQEAHAKL\DN
		i				LT\LMEIHTSG\TFLTQALNHMY
						KLRTNLQPLESTQSQDF
22911	53279	Α	23044	3	1038	TSSMAAGCSEAPRPAAG\SDGS
	l	l				LVGQAGVLPCLELPTYAAACA
						LVNSRYSCLVAGPHQRHIALSP
i		1				RYLNRKRTGIREQLDAELLRYS
						ESLLGVPIAYDNIKVVGELGDIY
ł						DDQGHIHLNIEADFVIFCPEPGQ
1	!				İ	KLMGIVNKVSSSHIGCLVHGCF
İ						NASIPKPE\QLSAEQWQTMEIN
						MGDELEFEVFRLDSDAAGVFCI
						RGKLNITSLQFKRSEVSEEVTEN
						GTEEAAKKPKKKKKKKDPETY
	1					EVDSGTTKLADDADDTPMEES
	1					ALQNTNNANGIWEEEPKKKKK
						KKKH\QEVSRTQDPCFSQGQLT
						S\SGYQSDHKKKKKKKKHSEEA
						EFTPPLKCSPKRKGKSNFL
22912	53280	В	23045	82	323	

3012

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22913	53281	Α	23046	3	1624	QLEIRAPTA DEIHVTVGEARNLI PMDPNGLSDPYVKLKLIPDPRN LTKQKTRTVKATLNPVWNETF
	l .	l				VFNLKPGDVERRLSVEVWDWD
	ŀ	l				
ļ	1			ŀ		RTSRNDFMGAMSFGVSELLKA PVDGWYKLLNQEEGEYYNVPV
		1				ADADNCSLLQKFEACNYPLEL
ŀ		1				YERVRMGPSSSPIPSPSPSPTDPK
į.	1	1				RCFFGASPGRLHISDFSFLMVLG
ŀ		l				KGSFGKVMLAERRGSDELYAI
ŀ		l				
		l l				KILKKDVIVQDDDVDCTLVEKR
•						VLALGGRGPGGRPHFLTQLHST
						FQTPDRLYFVMEYVTGGDLMY
ŀ		l				HIQQLGKFKEPHAAFYAAEIAI
	1					GLFFLHNQGIIYRDLKLDNVML DAEGHIKITDFGMCKENVFPGT
		l				TTRTFCGTPDYIAPEIIAYQPYG KSVDWWSFGVLLYEMLAGQPP
						FDGEDEEELFOAIMEOTVTYPK
		l				SLSREAVAICKGFLTKHPGEAP
	1			İ		GASGP*WGNLTIRAHGFFPLGF
		l				DWERLERLEIPASFSRPRPCGRS
						GENFDKFFTRAAPALTPPDRLV
l	1	l				LASIDQADFQGFTYVNPDFVHP
1		1		į.		DARSPTSPVPVPVM
22914	53282	Λ	23047	h	1522	MLRAPROATROHHFAHIDIVQV
22714	33202	ľ.	23047	1.	1322	TCFRYLRRSALAPAPAMPSPVR
		1				VDRGSFPPVAGGNHTRRRGGTP
		l l		i		OEGVTREALAFFDVVATTGAE
ł	i	1		l		MMWSNFFLQEENRRGAAGRR
İ		1		i		RAHGOGRSGLTPEREGKVKLA
	İ	1		i		LLLAAWAPRWRCCPWAPSSG
		1				WTQHLQGQRQRRVRSGPTWG
	i	l				CGRRAPSGCGRRTCPWTGTPA
		l				APRSCPEASERNSEHPALICTLP
		1		İ	İ	QPHMLKRRARRAPGSMAPPLP
		l		l		RPPOAASVPVFPPDWEPLESRA
	į	l				RSKSPLHSQHRPAQETGRHSRP
		l				RTSTRKROOKHSTEKATHTPER
		l				APKPQTRENNPKGRAQGSPAPR
		l				DHNPASRHQKTSNDKRSTHDT
	İ	l				HKTRKDSQTRRKKTRNNEKQQ
1	1	I		1	1	PDR/IGPQIIAYQP/YGKSVDWW
1	1	1			l	SFGVLLYE/MLAGOPP/FDGEDE
I	1	1				EELFQAIMEQ/TVTYPKSLSREA
I	1	1				
I	1	1		1		VAIC/KGCGRSGENFDK/FFTRA
I	1	1				APALTPPDRLV/LASIDQADFQG
1	1	I	1			FTYVNP/DFVHPDARSPTSPVPV
L	L		L	L	L	PASSPTRPQLLENKFGC

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amine acid	
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
22915	53283	A	23048	505	2345	I ISIWGMVMMNPTPSSITLRRMM
22713	33263	<u> </u> ^	23048	1303	2545	SLFPASLTTKYGGFYINSGTLQF
		l		1		RQASESEDDFIKEKKKKSPKKR
		1	1			KLKEGGEKIKKKKKKDDTYDKE
				1		KKSKKSKFSKAGFTALNASKEK
		1		1	l	KKKKYSGALSVKEMLKKFQKE
					ŀ	KEAQKKREEEHKPVAVPSAEA
ŀ		1				QGLRELEGASDPLLSLFGSTSD
		1		l	i .	NDLLQAATAMDSLTDLDLEHL
				1	1	LSESPEGSPFRDMDDGSDSLGV
		1		1		GLDQEFRQPSSLPEGLPAPLEKR
					l	VKELAQAARAAEGESRQKFFT
						QDINGILLDIEAQTRELSSQVRS
				ı		GVYAYLASFLPCSKDALLKRA
				i		RKLHLYEQGGRLKEPLQKLKE
İ		1				AIGRAMPEOMAKYODECOAHT
ŀ		Į.		i		OAKVAKMLEEEKDKEORDRIC
						SDEEEDEEKGGRRIMGPRKKFO
						WNDEIRELLCOVVKIKLESODL
				1	İ	ERNNKAQ\AWEDCVKGFLDAE
l	1	1		1		VKPLWPRGWMQARTLFKESRR
l		1		1		GHGHLTSILAKKKVMAPSKIKV
						KESSTKPDKKVSVPSGQIGGPIA
						LPSDHQTGGLSIGASSRELPSQA
	ŀ			1	ł	SGGLANPPPVNLEDSLDEDLIR
				1		NPASSVEAVSKELAALNSRAAG
				I		NSEFTLPAPSKAPAEKVGGVLC
		l		I		TEEKRNFAKPSPSAPPPASSLOS
22916	53284	A	23049	1	379	AYRNFTSASDVWSYGIVMWEV
220		1		ľ		MSYGERPYWDMSNQD\PNTAL
				1		LDPASPEFSAVVSVGDWLOAIK
				1		MDRYKDNFTAAGYTTLEAVVH
				l .		VNQEDLARIGITAITHQNKILSS
		İ				VQAMRTQMQQMHGRMVPV
22917	53285	Α	23050	l	123	
22918	53286	Α	23051	3	221	
22919	53287	С	23052	80	379	
22920	53288	A	23053	71	924	AFSVNSSNSWEKA*SQKCLQV
						NYLSLSSASYLP/CMTCPPSAPL
						NLISNVNETSVNLEWSSPQNTG
						GRQDISYNVVCKKCGAGDPSK
		1				CRPCGSGVHYTPQQNGLKTTK
						VSITDLLAHTNYTFEIWAVNGV
		1		1		SKYNPNPDQSVSVTVTTNQAAF
				1	1	SSIALVQAKEVTRYSVALAWLE
				1	1	PDRPNGVILEYEVKYYEKDQNE
				1		RSYRIVRTAARNTDIKGLNPLTS
	l			1	1	YVFHVRARTAAGYGDFSEPLE
				1	1	VTTNTGNKDHPDSVTSEAIFIIQ
1				1	1	LNPITVKSKELSVIFGSADILLQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22921	53289	Α	23054	1	1102	MRWLLKWERKYTLYSSGVLK
22921	33289	^	23034	[*	1102	HKNLLESLLGQFLGPPAQRFDS
İ						AGGVRPKNDGRFTVIQLVGML
		l				RGIGSGMKYLSDMSYVHRDLA
						ARNILVNSNLVCKVSDFGMSR
	ŀ					VLEDDPEAAYTTRGGKIPIRWT
	ŀ					APEAIAYRKFTSA\SDVWSYGIV
						MWE\VMSYGERPYWDMSNOD
						VIKAIEEGYRLPPPMDCPIALHO
1						LMLDCWOKERSDRPKFGQIVN
					İ	MLDKLIRNPNSLKRTGTESSRP
ļ	l					NTALLDPSSPEFSAVVSVGDWL
l						OAIKMDRYKDNFTAAGYTTLE
						AVVHVNQ/EVSTQRCNTKGDL
	İ				İ	ARIGITAITHQNKILS\SVQAMR
						TQMQQ\MHGRMVPRLSTVLNK
		ļ	20045	140	2220	LKTLEISLPHPCTLIEELHFFNFV
22922	53290	Α	23055	148	3229	NAFRFHLLNPAGESHDRLQDLL
ĺ						SAVRSPTKRQEQRWHRRTMAG
						IFYFALFSCLFGICDAVTGSRVY
						PANEVTLLDSRSVQGELGWIAS
						PLEGGWEEVSIMDEKNTPIRTY
		1				QVCNVMEPSQNNWLRTDWITR
		1				EGAQRVYIEIKFTLRDCNSLPG
l		ł	ŀ		1	VMGTCKETFNLYYYESDNDKE
1		1	Į.			RFIRENQFVKIDTIAADESFTQV
Ì						DIGDRIMKLNTEIRDVGPLSKK
		l	i i			GFYLAFQDVGACIALVSVRVFY
		_				KKCPLTVRNLAQFP
22923	53291	Α	23056	57	450	VGDCLTSRGMSWVQATLLAR/
1		l		1		ALCRAWGGT/CRGALTGTSISQ
1		l		1		VPRLAP\RGLHCSA/ASVSSEQS
1		l				LVPSPPEPRQR/PPGGERDKASF
1	l	l	l	İ	1	LQTVQKFADTSVVDSGHIDFIY
1		l		1		LALRKMREYGVERDLAVYNQL
1		l				LNIFP

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X-Unknown,
NO:	of peptide	hod	in USSN	location of first	endon for last amino acid	*=Stop eodon, /=:possible nucleotide
	sequence	1	09/540,217	eodon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
22924	53292	-	23057	89	11502	VGDCLTSRGMSWVQATLLARG
22924	33292	A	23037	109	1302	LCRAWGGTCGAAL/SOEPPSLR
						SLASLPRGLHCSAAAHSSEQV\L
					i	
		1				VPSPPEPRQEGPPRLLVPFEGPV LGOGAGLGERGOGRAFL/LDGO
						RNLRTTSVRKRGPH/VDLFYLA
						LRKMRE\YGVERG/DLAVYNQL
ſ						LNIFPKEVFRPRNIIQRIFVHYPR OOECGIAVLEOMENHGVMPNK
1						
						ETEFLLIQIFGRKSYPMLKLVRL
						KLWFPRFMNVNPFPVPRDLPQD
						PVELAMFGLRHMEPDLSARVTI
		l				YQVPLPKDSTGASKIPPKPHIVG
					ľ	HRACMSPHQESKKSRSSKAAL\
						ARHNPS\RPVF\VEGPFSLWLRN
					l	K\CVYY\HILRA\DLLPPEEEGK
					1	WKRTPEEWE\LYYP\RQLDLEY
					i	VRSG\WDNYEFDINEVEEGPVF
				1		AMCMAGAHDQ\ATM\VKWI\Q
į		1		1		GPAGDQPQTLAQNPPWSFRLRR
						VQPGELPDILLQGWRSP/HPAR
		_				GPPRKKTTTCKRQQQGPELV
22925	53293	A	23058	1	2517	HITTERIODOL AD ADOCDA UPOLI
22926	53294	Α	23059	1	927	IIVVEFVSRGLARAPSGPAVFSH
						LAMARLPLRGVRWGCLLTAVH
				1		PEPPTACREKQYLINSQCCSLCQ
1					1	PGQKLVSDCTEFTETECLPCGE
I						SEFLDTWNRETHCHQHKYCDP
1						NLGLRVQQKGTSETDTICTCEE
						GWHCTSEACESCVLHRSCSPGF
					i	GVKQIATGVSDTICEPCPVGFFS
						NVSSAFEKCHPWTSCETKDL\V
ĺ						VQQAGTNKTDVVCGPQDRLRA
						L\VVIPIIFGILFAILL\VLVFIKKV
						AKKPTNKAPHPK\Q\EPQEINFP
İ				1		DDLPG/SSNTAAPVAGRIFTWG
		_		L	100	GQPV\TQEDGQRESHFSCRERQ
22927	53295	В	23060	24	439	
22928	53296	A	23061	1	582	
22929	53297	С	23062	1 0 40	2607	+ TOUTTOE OF COUNTY OF THE COUNTY
22930	53298	Α	23063	248	430	ATNYTREQDSMWNCVR*GRTT
1		1		1	I	ATTHSVTQRTDERQRRADTAL
00001	50000	Ļ.	22064	200		TLADYRNTSVTGVFAGLF
2293 I	53299	A	23064	776 284	1426	VEDI I DOOVEEDOVEDOV 1 431
22932	53300	Α	23065	284	598	VFRLLRGQKSEPGKSRPKAL*N
1	1	1		1	I	KPANTPVQSHISLYAAGIINSDS
1		1		1		KVR*SCRQSSARQSGSYL*TSPR
	1	1		I	i	SPQNRQQTGEITATVIAFGRTRR
22224		L.	02055		2500	TAPDITPHPASAR
22933	53301	A	23066	3139	3792	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22934	53302	Ā	23067	2	830	GVPNMA APODVHVRICNOEIV
1		l				KFDLEVKALIODIRDCSGPLSAL
		l				TELNTKVKEKFOOLRHRIOPVL
		l				YORAFIWTASTFFFKLTYSLTDF
		1	ŀ			SSTOHDFNSPTTPVTFSDL\EQL
		l	ŀ			AKEODKESENOLLLOEVENHK
		1				KOMLSNOASWRKANLTCKIAI
						DNLEKAELLOGGDLLRORKTT
						KESLAQTSSTITESLMGISRMM
1		ı				AQQVQQSEEAMQSLVTSSRTIL
		1		ľ		DANEEFKSMSGTIQLGRKLITK
		1				YNR\RELTDKLLIFLALRLFLAT
						VLYIVKKRLFPFL
22935	53303	A	23068	1231	1458	
22936	53304	В	23069	1	2133	
22937	53305	Α	23070	722	957	VFRLLRGQKSEPGQKQAKSTLE
		i	1			QTSEYPSNRGIPIIRQSECSIRTPL
		l				PFVRALRD*VRGCGCAPLSSVD
		_				SFFCCTEAL
22938	53306	Α	23071	2	1137	VPNMAAPQDVHVRICNQEIVKF
						DL\EVKAPIQDIRD\CSGPLS\AL
						TELNPKVKEKFQQLRHRPTGP/
			l			VEQLAKEQDKESEKQLLLQ\EV
						ENHKK\QMLQQFRPSWEGKLN
						SSPGKICNSDNLEKAELLQGGD
						LLRQRKTTKE\SLAQTSSTIT\ES
						LMG/ISGRMMAQQVQQSEEAM
						QSLVTSSRTILDANEEFKSMSGT
			İ			\IQLGRKLITKYNRRELTDKLLIF
		1				LALALFLATV\LYIVKKRLFPIF
	l					VRFPKVPVLAPFNSLFSGICPWF
						PEAPRLPKAEAHTPLRFAPQLP
		l				CRLGWEHQCPPLFLQSSWLVK
	1					GEQSPWEEGPGIGMPPLGTYEP
						PPSHHCALWEAAAAWSTLKSE
						PPRTRMPLLSPAPIVPLNARSGG
		<u> </u>			***	GKREER
22939	53307	Α	23072	64	665	RVRGGEAELAMAAAEEE\SGAP
						NVPNRERGGAGRPKTPFECNIC
	1					LETAREAVVSVCGHLYCWPCL
						HQWLETR\PERQECP\VCKAGIS
	İ					REKVVP\LYGRGSQEAPRIPELK
						TPPPPPRGQR\PAPESRGGF\QPF
		1	1			GDTGGFHF\SFGVGAFPF\GFFT
		1				TVF\NAHEPFRRGTGVDLG\QG
		L		L	L	HPSLQLGKEFPLPGFSAIFFLFW

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nuclcotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22940	53308	A	23073	1	416	MQRRRRRRLLLLPPRRQTDRAL PPPPSPGSMPPPLPPPP/PPARGG GLRPGDGLYPK/PSPARRGSGA
		1		l		RTSWCTMCAPPSTSAPRASRRP
		1		l		RPSSCATRPPTSKPPPA\GHAPH
		l				PPPRDLGGGLDSGVQSDGVLQ
				ŀ	1	HLORPGH
22941	53309	Α	23074	1	1713	
22942	53310	Α	23075	1305	1538	LKVKRNCTSS*SPTIGHD*QSNL
						SLPPSDFAKPPFPFSLLRCSKFR
						KLKAAYRKRKKGHKSSLSLSV
		1				KINKTAAANK
22943	53311	Α	23076	49	355	
22944	53312	В	23077	253	471	
22945	53313	A	23078	1267	1497	LKVKRNCTSS*SPTIGHD*QSNL
						SLPPSDFAKPPFPFSLLRCSKFR
İ						KLKAAYRKRKKGHKSSLSLSV
		l				KINKTAAANK
22946	53314	Α	23079	3	1237	HEVAGAVPGAIMDEDYYGSAA
		1			1	EWGDEADGGQQEDDSGEGED
1					ŀ	DAEVQQECLHKFSTRDYIMEPS
		l				IFNTLKRYFQAGGSPENVIQLLS
		i			ł	ENYTAVAQ\AVNLLAE\WLIQT
1		1				GVEPVQVQDTVEYHLKSLLIKH
						FDPRIADSMFTEEGETPAWLE\Q
				i		MIAHTTWRDLFYKLAEAHPDC
						LMLNFTVKLISDAGYQGEITSV
l						STACQQLEVFSRVLRTSLATILD
ŀ						GGEENLEKNLPEFAKMVCHGE
1						HTYLFAQAMMSVLAQEEQGGS
}						AVRRIDQEVQRFAQEKTVLTQE
		1				AIITVKGVSLSSYLEGLMASTIS
						SNASKTIVAAMSNLVPPVELAN
						PENQFRVDYILSVMNVPDFDFP
						PEFYEHAKALWEDEGVRACYE
						RSNEYQLIDCAQYIFVLEMQVP
		l	L			CCEEAQSHPGQHGGVQVGRN
22947	53315	A	23080	1358	1589	LKVKRNCTSS*SPTIGHD*QSNL
1		1		1		SLPPSDFAKPPFPFSLLRCSKFR
		l	İ			KLKAAYRKRKKGHKSSLSLSV
	L					KINKTAAANK
22948	53316	A	23081	33	458	GITGSTDASGVTMKFNPFVFSD
						RRKNRKRHVNAPSHVRRKIMS
			ĺ	l		SPLSKELRQKYNVRSMPIRKDD
	l			1		EVQVVRGHYKGQQIGKVVHV
1	1	1				YRKKYVIYIERVQREKANG/DK
			ĺ			DRKKILERKAKSRQVGKEKGK
L	L	<u>L</u>				YKEELIGKMQEYIEP
22949	53317	Α	23082	1	576	
22950	53318	В	23083	37	276	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22951	53319	Α	23084	124	549	NNSLSSPVQADVYFPRLIVPFCG
						HIKGGMRPGKKVGVMGIVDLN
	1	l				PKSFAINLTCGDSEDPPADGAIE
	l	ĺ				LKAVFTDRQLLRNSCISGERGE
		1				EQVSNPLLSIHSTTMPFRVEILC
		1				EH\HCFR\VFVDGHQLFDFYHPH
		1				SNRYLAI
22952	53320	Α	23085	1	215	QGFIKPPLKEENFTLPPPGKFGP
		1				PKGSLNSPPS/PSSP*SPGGGILLL
		l				VYYLYRALFQKGCKEAAWLEY
						PLT
22953	53321	Α	23086	533	983	PWGPGCNKRWPCLAKDPPPTP
		1				QSSGVGQPFGGAEPAGAYIRTG
		1	İ			PHLGPPPPPPPDSSTNLHSPNPS
	İ	ı				DDGADTPLAQSDEEEERGDGG
	1					AEP\G\PAASSGPLAYRPDPAGY
		l				SPTSRPPGPSPEPLG/AEDPDSLL
	1					ASRAPRGGNGWWDCGTLS
22954	53322	Α	23087	1422	1670	NRYIDKTGWAWWLTPVIRHFR
	ŀ	1				RPRWVDCLRLGVREQPGQHSKI
						KSLQKNAK\ISQAWWHAPVDP
		l				ATWKADAGGSLEPGSLRLQ
22955	53323	Α	23088	2	663	VLLDERSAALDGAKRDGTLAL
			i			AAGALCREARAAQVFFLKGGY
						EAFSASCPELCSKQ/INVSANCP
		1				NHFEGHYQYKSILCGMTTHKA
		1				DISSWFNEAIDFIDSIKNAGGRV
						FVHCQAGISRSATICLAYLMRT
						NRVKLDEAFEFVKQRRSIISPNF
		1		i		SFMGQLLQLESQVLAPHCSAEA
						GSPAMAVLDRGTSTTTVFNFPV
				1		SIPDHSTNSALSYLQSLITTSSHC
22956	53324	Α	23089	3	206	

3019

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22957	53325	Α	23090	982	2252	LFLEKRRPALGPENRQRAPQGR
						STLGALRPAMVMEVGTL\DAG
		İ				GLRAL\LGERAAQLPACLDCRS
						FF\AFNAGHIAGSVNVRFSTIVR\
	ł					RRAKG\AIGALE\HIVPNAELR\G
						RLLAGAYHAVVLLDERSAG\LD
	İ	i				GAKRDGTLALAAGRA/LCREA
		i				RAAQVFF/LSKGGYEA\FSASCP\
						ELCSKQSTPMGLS\LSLSTSVPD
		l			1	SAESGC\SSCSTP\LYD\QGGPVE
	ŀ	ı				ILPFLYLGSAYHASRKD\MLDA
		l				LGHRLPWINVLSQIVPNHF\EGH
						\YQYKSIPVEDNHKADISFWFN\
	İ					EAIDFIDSIKNAGGRVFVHCQA
		l				GISRSAT\ICLAYLMRTNRVKLD
	i i					EA\FEFVKQRRSIISP\NFQLHGA
		ŀ				SLLQFEFPRMLGPALFRAEAGK
						PRQMAVL\DRGT\STTTVFNFPV
		l				SIPVHSTNSALSVPFRAPLTDLS
	1	1				QLVERPTGR
22958	53326	Α	23091	13	287	
22959	53327	Α	23092	1	492	
22960	53328	Α	23093	1	636	
22961	53329	A	23094	1	506	PTRTWTRGRIPRLSAQPSRGAR
		l				GTMADPRVRQIKIKTGV\VKRL
		ŀ		1		V\KEKVLYEKEPKQ\QEEKIEKM
		l				RAEDGENYDIKKQAEIL\QES\R
		1	1			MMIP\D\CQRR\LEAAYLDLQRI
i						LENEKDL/EKEAEEYKEAR\LVL
	Į.	l				DSVKLESLKLFSYGVVFALKPR
		<del>  -</del>				GPILQSIIFDHVLCVQV
22962	53330	A	23095	1	786	ULC A FUNCCI PROPET A FEBUR
22963	53331	Α	23096	183	1059	VLGAFIWQGLDPSPETAFTDHR
ļ.		1		1		ATEWGTNDQSLCLNSP*\DAPS
		l				AKHEYQASVSGSVSSKPEIDLQ
		l				LYDEVRDQDVPVKRNFARIVV
		l		1		NVSDTNDHAPWFTASSYKGRV
	ŀ	l				YESAAVGSVVLQVTALDKDKG
	İ	l				KNAEVLYSIESGSHPPYKTAAD
		l				VLLKAPPPGWRPTNTKPVHSTE
		l		1		AQPTQNQRTQQKHNQGASRSP
	1	l	1	1		RHSFATSTRAGAGICGCKTRRW
		l		j		MTRWMEESRGSLQTLPSTRLEP
		1				GSSAAWLDPEEQAQSLQFSSQE
				1		APFLGEGGENHIKGVPHGTKES OOOPLNHRSSL
22964	53332	Α-	23097	1	1224	QQQFLININGSL
22964	53333	_	23097	1	660	
22960	22333	Α	23098	Li.	1000	l

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
22966	53334	Α	23099	129	595	LFPRLLSCLTTPPHCSFSICFVIC
1						SRTLILKGSSLMYVFCLPNTAIV
						MALSPRGWRSKFGMPWHSKPT
1	l	ł				WIMTL*FPLEKTGALSLTSDTS
		l				MFTVVVLTLGGLPLSEA*T/GG
						NYDSHFDVDKGTGTIIVAKPLD
						AEQKSNYNLTVEATDGTTTILT
22967	53335	Α	23100	1	1782	MAARIEVIHELSIRLLLMKTDL
						DAATAALLNVQPAKSGDNTSG
						APFPGKISQTLLKDPHNGYHVV
		1				FALAGSPKDADDTSIYMFYQK
	İ					VGDNSIDSWKNAGRVFKDSDK
		l				FDANDPILKDQTQEWSGSATFT
						SDGKIRLFYTDYSGKHYGKQSL
		l				TTAQVNVSKSDDTLKINGVED
	İ					HKTIFDGDGKTYQNVQQFIDEG
						NYTSGDNHTLRDPHYVEDKGH
						KYLVFEANTGTENGYQGEESLI
						NKAYYGGGTNFFRKESQKLQQ
						SAKKRDAELANGALGIIELNND
						YTFEKSNEAADHFKHGGGATIK
					İ	GRVADNGKPQLSSLTYIDIRVIE
						ESIYPPAILPLEIFITASGEEYSGO
						VIGKIHATDQDVYDTLTYSLDP
						QMDNLFSVSSTGGKLIAHKKLE
						IGQYLLNVSVTDGKFTTVADIT
						VHIRQVTQEMLNHTIAIRFANL
			1			TPEEFVGDYWRNFQRALR\NIL
						GVRRNDIQIVSLQSSEPHPHLDV
1						LLFVEKPGSAQISTKQLLHKINS
1			1			SVTDIEEIIGVRILNVFQKLCAG
1						LDCPWKFCDEKVSVDESVMST
1		l				HSTARLSFVTPRHHRAARQQ\R
						GPPLHNPADT*TTARKYNCDKR
22968	53336	Α	23101	184	13958	AMGRHLALLLLLLLLFQHFGDS
		1				DGSQRLEQTPLQFTHLEYNVTV
j		1				QENSAAKTYVGHPVKMGVYIT
j		l				HPA WEVRYKIVSGDSENLFKAE
		1				EYILGDFCFLRIRTKGGNTAILN
		1				REVKDHYTLIVKALEKNTNVE
1	1	1				ARTKVRVQVLDTNDLRPLFSPT
1		1				SYSVSLPENTAIRTSIARVSATD
1		ı	ĺ	1	I	ADIGTNGEFYYSFKDRTDMFAI
	l	1	1			HPTSGVIVLTGRLDYLETKLYE
	l	1		1		MEILAADRGMKLYGSSGISSMA
	l	1	1			KLTVHIEQANECA
22969	53337	С	23102	56	331	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22970	53338	A	23103	2249	2680	GEFLKTREFSIYFLEKYTTFPGE NLEHGFWDSSTSPPVLLHPGIPA SRI/PVSSIFTXVPPKKSATKP/PV SIWGAPPLGPPGFWGCCPTGEQ RGVCHPSEPHPPVVDFRKKPPF LCVSLSLINIFKSNKFLLTTCNL VIFYTCSL
22971	53339	A	23104	739	1672	COEAASEFGCPLHTPAMFLIRIL GGWLPRPWGRRKPMRPDPPYP EPRRVDSSSENSGSDWDSAPET MEDVGHPKTKDSGALRVSRAA SEPSKEPEPQVEQLGSKRMDSLK WDQPISSTQESGRLEAGGASPK LRWDHVDSGGTRRPGVSPEGG LKGVPGFGAPLEKPGRBEKLLG WLRGEFGAPSRYLGGFEECLG WLRGEFGAPSRYLGGFEECLGS STINLTHLELELLASALLAGLELASAL LSFLAALIGHAVLSLTTAHL LFSLAALIGHAVLSLTTAHL HFACLFGLLQALVLAVSLREPN GDEAATDWESEGLEREGEEQR GDPGKGL

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SEQ ID	SEQ ID NO:					Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	or pepude sequence	detection, v-possible nucleotide insertion)
	<u> </u>	<u> </u>				
22972	53340	Α	23105	122	2437	RLELKLPLWTMRFTEMELLGH
				ĺ		LSIFKITGKQKQLTRTAPGPSGQ
	ł			l		LRSNQTTGNSLEPSGQPRSSQTT
			1			GNSLEPSGQPRSSQTTGNSLEPS
			i			GQPRSSQTTGNSLEPSGQPRSSQ
		1				TTGNSLEPSGQPRSSQTTGNSLE
	Ì	1				PSGQPRSSQTTGDSLEPSGQLRS
		1				NQTTGDSVQPNNREQPGTVRS
		1				APLQPNNREQPGTVRSAPLQPN
l		1				NRGQRPTKQQGTTWNRQVSSA
		1		1		PTKQQGTAR\PSGQPRSNQTTG
1	1	1				NSLEPSSQLRSNQTTGDSVQPN
İ						NREQPGTVRSAPLQPNNRGQRP
				1		TKQQGTAR\PSGQPCSNQTTGN
						SLEPSSQLRSNQTTGDSVQPNN
						REQPGTIRSAPLQPNNRGQRWD
						RQVSSAPTKQGTALGPSG\SSAP
		1				TKQQGTALEPSSQLRSNQTTGN
l	1			1		SLEPSGQPRSNQTTGNSREPSSQ
			i .			LCSNQTIGNSQDRQVSSAP/IQT
1	1	1				TGNSQDRQVSPAPTKQQGTAG
		1		ļ		NRQVSST/PIQTGNSVQPNNREQ
}						PGTVSGAM/VWVATPVKGPAR
				1		VETSAHKVSAAGLPKKKT/QPG
	1	1	1	1		VTGGAPRLTLNTVLPKHSRTPS
		1		i		RHEAGSTQALYHNGQAQYALS
		1				FPHEAAPMAAFYVVTGIEVTTA
		1				LTLEEAPCTVEITTALTLEEPPC
						TVFWNKTERNTRTLVQDKGSN
			ļ			TKGFVQTATCCFRADCKASRPS
		1				PHPPSPHREWTSDLRHKQVSKF
		_				PGAHRLPGHPELPIQEAWDRGE
22973	53341	Α	23106	605	813	LFVFINFRENPALLRWAYAR\TI
						NVYP\NFRPTPKNSLMGALCGF
		_				GPLIFIYYIIKTERVSIQTRCLVF
22974	53342	Α	23108	273	936	AYSNKVPSWNQKA\PSPTKPAG
1						TLIFDFPAVKTVRNKFVFFINYP
					1	VSGYTEDLKKFLKPYTLEEQKN
						LTVCPDGALFEQKGPVYVACQ
						SPISLLQACRGMNDLDFGYSQR
1		1				NPCILVKMNRIIGLKSEGVPRID
		1	l			CVSKNEDIPNVAVYPHNGIIDL
						KYFPYYGEKLHVGYLQPLVAV
1		1	1	1		QVSFAPNDTGKEVTVECKIDGS
		_				ANLKSQDDRDKFLGRVMFKIT
22975	53343	Α	23109	104	478	ETEVGGLVEASMSHNCTTALQ
		1	1	1		PGQHSQKENLQNKPIPGREKY*
		1		1	1	VSIEFELIYFP/YYCQVGYLQPL
1		1	1	1	1	VAVQVSFAPNNTGKEVTVECKI
		1		1		\DGSANLKSQ\DDRDKFLG\RV
L			L		L	MFQNHSTVHSMSRVSSTE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22976	53344	Α	23110	3	903	LLSPVRAHTMTKNEKKSLNQSL
						AE\WKLFIYNPTTGEFLGRTAAS
		l				WGLILLFYLVFYGFL\AALFSFT
						MWVMLQTLNDEVPKYRDQIPS
						PGLMVFPKP\VTALEYTFSRSDP
						TSYAGYIEDLKKFLKPYT\LEEQ
				ļ		KNLTV\CPDG\ALFEQKGPVYV
			İ			ACQFP\ISLLQACSGMNDPDFG
			i		l	YSQGNPCILVKLNRIIG\LKPEG
					l	VPRIDLCFKRMKDIPNVAVYS\
						HNGMIDLKYFPYYGKKTAMF/
						VGIYQPIGLLVQVSFAP*QPLGK
						EVTVE/CAKIDGSSQPKKVQDD\
						RDKVFGDEFMFQIHSTVH
22977	53345	Α	23111	467	931	SRDSRRGTPLAGGRGRDFAGQ
						MGHRVQGRHGSRGRLPSGDAR
				ŀ		VNAERAEPVGAGAGGPGGSS
		ł				VRGFCGRSWSEA/RGRRRAGG
						GVH\IEP\RYRQFP\QLTQIPRCSQ
					1	SEF\FSGLMWFWIF\WRFWHDP
						K\KVLGHFPV/YPDPSQWTD\EE
						L\GIPPDD
22978	53346	Α	23112	1	390	VTIPTSLAELQGLPHIRVFLDNN
					l	PWVCDCHMADMVTWLKETEV
						VQGKDRLTCAYPEKMRNRVLL
		1			1	ELNSADLDCDPILPPSLQTSYVF
		1			1	LGIG*ALIGAIFLHYRNNGPLTIS
		1			1	SEGGLASLLCKDSRGVPLL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
22979	53347	Α	23113	3	1653	PQRIQRGVPTRGEEVAPGRRQE
		1				EEREQERGAERPDPPCVLSGGK
l		l				GRGNRPLREAPGGEGVSQVPA
						AAPLPRFHERKVFFFQTLPPAR
						ALRAQPPEPSERAPSQPSSGETR
l				1		AAMPGGCSRGPAAGDGRLRLA
						RLALVLLGWVSSSSPTSSASSFS
l		l				SSAPFLASAVSAQPPLPDQCPAL
		l				/CACRSEAARTVKCV\NRNLTE
l						VPTDLPAYVRNLFLTGNQLAVL
						PAGAFARRPPLAELAALNLSGS
						RLDEVRAGAFEHLPSLRQLDLS
			ŀ			HNPLADLSPFAFSGSNASVSAPS
						PLVELILNHIVPPEDERQNRSFE
						GMVVAALLAGRALQGLRRLEL
			Ì		ĺ	ASNHFLYLPRDVLAQLPSLRHL
		ļ				DLSNNSLVSLTYVSFRNLTHLE
ł						SLHLEDNALKVLHNGTLAELQ
						GLPHIRVFLDNNPWVCDCHMA
						DMVTWLKETEVVQGKDRLTC
				1	1	AYPEKMRNRVLLELNSADLDC
		1				DPILPPSLQTSYVFLGIVLALIGA
						IFLLVLYLNRKGIKKWMHNIRD
					1	ACRDHMEGYHYRYEINADPRL
						TNLSSNSGCLEEILEDRPRTTLH
22980	53348	A	23114	288	877	ASRLSLPTRTSMRSSSSTSPSSSS
		l				SSSSLCRFLLNSRVTDAAFNFLL
						VWYYCTLTIRESILINNGSRWG
						RAGPEGGEHGMGCQGPSGKGR
				l		EQGAGG*WWVFCSRV\KGWW
						VFHHYVSTFLSGVMLTWPDGL
						MYQKFRNQFLSFSMYQSFVQF
						LQYYYQSGCLYRLRALGERHT
			ĺ			MDLTVGESGVALASGHSPFLPS

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SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22981	53349	Α	23115	1	1273	MAPLHSILGNRVGLCQASVRA
						CPGATVSSQCRASLRSRLAWRG
	ł				1	SELPANRGRTWPGPGGLRLOA
	l					APRLAYLGRATLLVLTPYYDLP
					ĺ	PNVRFTSMPASPHTYFLVLGAPI
					i	
	1			İ		FLLLRSTWSYYHACRFEYATPN
						DRNTTYLAISGPTRRTYDVATF
						AYKDEYEKFKLYLTIILILISFTC
		l				RFLLNSRVTDAAFNFLLVWYY
						CTLTIRESILINNGSRPDGLMYQ
	1		i			KFRNQFLSFSMYQSFVQFL\QY
	1					YYQSGCL\YRLRALG\ERHTM\D
						LTVEGFOSW/MWRGLTFLLPFL
						FLW/YNSGQLFNALTLFKPGPR
						DPQLQGVGRCLCCGFPFLL\LFL
	l					
						GNFFTTLRVVHHKFHSQRHGS
	ļ					KKDLRLGLSPCRPRRGFLVPVC
						FWEGDWEGAPSSVRVSGGLFY
						SPLGFMGRCGPGKEDPGPSCPQ
22982	53350	Α	23116	238	398	
22983	53351	A	23117	1	386	
22984	53352	Α	23118	13	580	
22985	53353	В	23119	39	1171	
22986	53354	A	23120	381	795	PPSSKAPSSKGPAKIVCIYWALI
						LOSNEVTITEDKFNNLIKAAAV
		l				TVEPFWPSFFAKALASVNIGSLL
						CNAGVG/SMAPAACTAQAGGP
ı						ATSTTAASASFRNHNPRIAPKFR
						GVLAHEQYLSERATWVWAPNP
		_				NPKSLF
22987	53355	Α	23121	1	960	
22988	53356	A	23122	2	3952	
22989	53357	Α	23123	1058	1494	KSVRLVVN/YLRTQKAVVRVSP
	1	ı	1			EVPLQNILPVICAKCEVSPEHVV
1		l	l			LLRDNIAGEELELSKSLNELGIK
		1	l			ELYAWDNRRVPSAVPMOGPWP
		1				AEHFLOVNEOPPAGSPSPCPSSS
1		ı				SSSSGLRGGMOTANGPACALTP
1		l	l			
22222	50050	ļ. —	00101	105	200	TCAQTDCPCPTSH
22990	53358	Α	23124	195	398	TDSSSALQPQHRGRPALRQRTP
						PRRSP/SPGLSPPRGASPAS\SGA
1		1	l			PARGVAFGGDEPPIGAEAMYSK
			l			P
22991	53359	Α	23125	2	155	
22992	53360	Α	23126	1	339	
		<del></del>				

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22993	53361	A	23127	150	1017	LGRWRODOVPHGLECRSOSEW
22993	33301	l^	23127	150	1017	SCTDSROHSSS\YSDSRROEIGK
1					1	MRYVSVRDFKGKVLIDIREYW
						MDLKRLPRKPRTGEAGSGSLSS
				ŀ		PGADTPVEAAVTWRLGLGSVD
						VGSFRAWQRGKSAAVGHTVSP
				ŀ		RWPRWQGGDSSALSMASLSTL
					ŀ	TCSWGLRYSALSPTTSAQREEG
						SALYRHLSPPSLRFYPTVKWAT
					ľ	LAITRPRPPAPRSHSHHPQPRAL
						CPVWVLFQHPKTHGENLGDRW
						KEGSALVSSKGTPAKLSSGRIR
1				ŀ		NALQNTLGEVSVARIQTHRGIF
						VETTEARL
22994	53362	A	23128	I	584	APIECGGIPSLPVLCQSRANDQE
ł		l				GVRLLEPSEAMPKSKELVSSSSS
						GSDSDSEVDKKLKRKKASLL/P
		1				ENPVKK\QKTGETS\RALSSF*TE
1						PASSRDDNMFQIGKM\RYVSV\
		1				RDFKGKSA*FDIREYWMDPEGE
						MKPERKGISLNP\QQWEPSLKE
		l				QISGPLIDASKKTCKISEPILIKPC
		ı				TVPVVLICLVTVAVCF
22995	53363	Α	23129	186	687	LLWARGLGRAKSAVPTVSTML
į		1				GLPWKGGLSWALLLLLLGSQIL
ļ		l				LIYAWHFHEQRDCDEHNVMAR
İ						YLPATVEFAVHTFNQQSKDYY
ĺ						AYRLGHIFNSWKEQVESKTV\F
j						SMELLLGRTRVWEYLKTTIDNC
						HFOESTELNNTFTCFFTISTRPW
ĺ		ı				MTOFSLLNKTCLEGFH
22996	53364	A	23130	232	402	LFKAWECFFLIFLOVVITRLKLD
		1				KDRKKILERKAKSRQVGKEKG
		ı				KYKEETI\EKMOE
22997	53365	A	23131	637	758	NVEEKNRKKILERKAKSCOI/K
						KEKGEYKEETIGTMRECGL
L	1			L	L	TELECOL TELECTION TO THE COLD

SEO ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleatide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	coden for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
						L
22998	53366	Α	23132	1	1095	IVFLLYLETCLEVMDDKPNPEA
		l		l	1	LSDSSERLFSFGVIADVQFADLE
			i	l		DGFNFQGTRRRYYRHSLLHLO
		1	1	i		GAIEDWNNESSMPCCVLOLGDI
		1		1		IDGYNAQYNASKKSLELVLW\Q
		l		į.		FKRRKGPVHHTWGNHEFYNFS
		l				REYLTHSKLNTKFLEDQIVHHP
				l .		ETMPSEDYYAYHFVPFPKFWFI
		l				LLDAYDLS/ALGAWDQSSPKYG
		1		İ		GSVWKILEGSHNP\NTGNWNSS
1		l				SRDFSEA/RSLFQFNGRVSAQEQ
1		l				
		l				LNWLNEVLTFSDTNQEKVVIVS
		l		i		HLP\LYPGPPFDNVCLAWNYRD
		l	ļ.			ALAVIWSHECV\GVSLLGHTHD
						GGYSEDPFGVYHG\NLEGVIET
ŀ		ł				APDSQRFGTVHVYPDKMMLKG
						RGRVPDRIYELQGKEEPSHC
22999	53367	С	23133	235	382	
23000	53368	В	23134	63	293	
23001	53369	Α	23135	160	330	
23002	53370	В	23136	1	870	
23003	53371	Α	23137	2	1013	
23004	53372	Α	23138	I	1552	
23005	53373	В	23139	120	1034	
23006	53374	Α	23140	1	584	RTRGRRSLAFLLLAERGAAAA
		l				VLWSGCRRFRARLGCLPGGLR
İ		l				VLVQTGHRSLTSCIDPSMGLNE
ľ		ı	1			EQKEFQNAAFDFAAREMAPNM
1		l				AEWDOKELFPVDVMRKAAOL
ŧ		l				GFGGVYIQTDVGGSGHVRLDT
1		l				SVIFEALA\TGCTSTTAY\IRHPO
1		ı				CPOSOHSFVOEGGTHYKAGHL
1		l				CCROWGPRACLPPGRPCHRGA
23007	53375	A	23141	2	412	CONQ II OTTUTODITI ONU CITACON
23008	53376	Ā	23142	47	487	
23009	53377	A	23143	77	459	GWKGIEHTLGRCLEAOGVINY
2300)	33377	ľ.	23143	l''	1737	RMLLSWRRWLVAHPVF*SFSCE
1	i					VGPVGQS*RMTMLHCRHSDPV
l						AKRLSCRRGDFASVVVLCPPPO
1						RLRTLKHFLFFGGLPLSIDFVMT
	44470	١		ļ.———	***	VDFCCHLHLTDFSSNFGF
23010	53378	A	23144	1	304	KVAGSPSPTPAGQQVLDVNSCL
			1			DSSVLDSSFLTFSGLHAE/AEGS
1		l	l			TGGPAALAYLGLKTLCTSQQAF
1		l	l	1		AGQMKSDLFLDDSKSDLEEPW
						ELRKEGAPRSRDRE
23011	53379	Α	23145	1	207	
23012	53380	Α	23146	1	3009	
23013	53381	A	23147	3	298	
23014	53382	Α	23148	1	636	
23015	53383	Α	23149	1	795	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23016	53384	Α	23150	125	1527	SSPROOTOLARVHP/DGRAO/LA
						PSCLPGLGLCPLPVCPPVCNCPA
						CPCHEHCGEAP*GVA\AKRGAL
						RVHPHPSCGGPRAMTVTLHGQ
		ŀ			•	DVLMLPGDLVGLQHDAGPGAL
						LHCSPAPGHPGPOAPYLSANAS
		ŀ				SWLPHLPAQLKGTWACPACAL
						RLLAATEQLTVLLGLRPNLGLR
						LPGRYEVRAEVANGVSRHNLS
İ		l				CSFDVVSPVAGLRVIYPAPRDG
						RLYVPTNGSALVLOVDSGANA
		ĺ				TATARWPGGSVSARFENVCPA
						LVATEVPGCPWETNDTLFSVVA
						LPWLSEGEHVVDVVVENSASR
1						ANLSLRVTAEEPICGLRATPSPE
					İ	ARVLQGVLVVLLAGSSGYLVG
						FKFLESHGSDSGSANSFHCLISQ
ŀ						NEFKTLLPDLTRAPRYSPVVEA
İ						GSDMVFRWTINDKQSLTFQNV
					l	VFNVIYQSAAVFKLSLSADSAP
						RPRWRLQSPRVGRLSPDPGFFG
						SPGPCQPRLQAFLI
23017	53385	A	23151	648	1387	ASTOSLLAWCPAWLSIPSTWPS
23017	33363	^	23131	048	1307	SFSSGCPG/CKAOPVYEVSRSSY
						VYLEGRCLNCSSGSKRGGYTFT
						LT\VLGRSGEEGCASIPLSPNRP
				l		PLGGSCRLFPLGAVHALTTKVH
1				l		FECMGWHDAEDAGAPLVYAL
1						LLORCROGHCEEFCVYKGSLSG
						YGAVLPPGFRPQFEVGLAVVV
						QDQLGAAVVALNRSLAITLPEP
						NGSAMGLTVWLHGLTASVLPG
				l		LLRQADPQLVIEYSLALVTVLN
						ETYTESL
23018	53386	A	23152	2	654	DKHIWLSIWDRPPRSCFTRIQRA
23018	33300	^	23132	<b> </b>	054	TCCVLLICLFLGANAVWYGAV
			1	I	l	GDSAYSTGHVSRLSPLSVDTV\
1				į.	l	AVGLVSSVVVYPVYLAILFLFW
				I		MSRSKV\AGSPSPTPAGOOVLDI
				l		DSCLDSSVLDSSFLTFSGLHAEV
1					i	INTLADHQHRGTDFGGSPSVLII
	-			1		TVSLRSYKFAISLCTTYLWVINT
				1		LADHRHRGTDFGGSPWLLIITV
	L		L		L	FLRSYKFAISLCTTYLC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
		1		sequence		
23019	53387	A	23153	2	4072	SRDPAGSYHLNLSSHFRWSALQ
23019	33361	l^	23133	ľ	4072	VSVGLYTSLCOYFSEEDMVWR
		l			\	TEGLLPLEETSPRQAVCLTRHL
		l			ì	TAFGASLFVPPSHVRFVFPEPTA
		1				DVNYIVMLTCAVCLVTYMVM
		l				
						AAILHKLDQLDASRGRAIPFCG
		1				QRGRFKYEILVKTGWGRGSGT
			1			TAHVGIMLYGVDSRSGHRHLD
		1				GDRAFHRNSLDIFRIATPH\SLG
						SVWKIRV\WHDNKGLSLALGFL
						Q\HVIVRDLQTA\QSTFFLVNDW
		<u> </u>				LSVETEANGGLVEKE
23020	53388	Α	23154	1	10922	MPPAAPARLALALGLGLWLGA
		1				LAGGPGRGCGPCEPPCLCGPAP
					Ì	GAACRVNCSGRGLRTLGPALRI
						PADATALTAMAKADGDVKKY
1						GSKRGALALTHGHSLLRDVSH
ł			1			NLLRALDVGLLANLSALAELDI
1						SNNKISTLEEGIFANLFNLSEINL
			1			SGNPFECDCGLAWLPRWAEEQ
			1			QVRVVQPEAATCAGPGSLAGQ
						PLLGIPLLDSGCGATLVGPHGPL
						ASGQLAAFHIAAPLPVTATRWD
						FGDGSAEVDAAGPAAS
23021	53389	A	23155	3887	14184	HVLCAWVTYMVMAAILHKLD
					ŀ	QLDASRGCAIPFCGQRGRFKYE
						ILVKTGWGRGSGTTAHVGIHA
				1		VWGGQPERPPAPGTATEPSTAT
						VWDIFQIRHPAQPGVACWKIRV
						FGTTTKGSALPRFPCSTSSSGTC
			- 4			RRHAAPSSWSMTGFRWRRRPT
						GAWWRRRCWPRVRPRSVFPLR
						HAALLRFRRLLGAELQRGFFDK
						HIWLSIWDRPPRSYFTHIQRATC
					Į.	CVLLICLFLGANAVWYGAVGD
					1	SAYSTGRVSRLNPLSVY
23022	53390	Α	23156	1	442	GGASGMGGIT\ASTVNQ\SLLSP
	1	1			1	LVLEVDP\NIQAVRTQEKEQIK\
1			1			TLNNKFASFIDKVRFLEQQNKM
1	1		1	l	l	LETKWSLLQQQKTARSNMDN
1	1		1			MFESYINNLRRQLETLGQEKLK
						LEAEFTTWGPPLPMPPATKQFN
1						CFFFLVQNKTSASSANVK
23023	5339I	A	23157	I	497	,
23024	53392	C	23158	208	422	
		_			·	L

NO:	SEO ID	SEO ID NO.	Met	SEO ID NO-	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
23025   53393   A   23159   81   1339   ELSSPRSSFSAPSRISAWFGPF   TPASTMSIRGTQKSYKGSTSG RAFKQAAS*QRSGPGFRISSS SRVGSNFRGGLGGGGGGG A MGGITAVTVNQ)SRALLSPL   EVDPNIQAVROPECEQIKTJ   KFASIFIDEVGFLEQONKML KWSLLDQQKTAKRSIMDING ESYINNLRRQLETLAQEKLK   AELGINMQGLVEDFKNNYED   NKSTETENEFYLIKKDVDEA   NEXTETENEFYLIKKDVDA   NEXTET							
23025   53393   A   23159   B   1339   ELSSPRSSFAPSRISAWFGFF   TPASTMSIRGTQKSYKGSTX   RAFKQAAS*QRSGPGFRISSS   SRVGSNNRGGLGGYGYGGA   MGGITAVTVNQISRALLSPL   EVDPNIQAVRTQEKEQIKTLI   KFASIFIDEVGFLEQQNKML.   KWSLLDQQKTAVRSNMDIN   ESYINNLRRQLETAQEKKLI   AELGNMQGLVEDFKNNYED   NKSTETENEFVLIKKDVEA   NELYEEIRELGGISDTSVV   DNSRSLDTDSIJAEVKTEISEN   RNISWLQAETEGLAGGRISLI   GIADAEQRRELVIKDANAKL   LEAALQQTKKLLEMENMSII   KTTSGVAGGVSLAVGGLTG   LSYGLSSSSRSGAGSSSFSRT   RAMVVKKIKTROCKLLTLS   LSYGLSSSSRSGAGSSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLTLS   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLT   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLT   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLT   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLT   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKLT   LSYGLSSSSRSGAGSSFSRT   RAMVVKKITROCKT   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKT   LSYGLSSSSRSGAGSSFSRT   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKT   LSYGLSSSSRSGAGSSFSRT   LSYGLSSSSRSGAGSSFSRT   LSYGLSSSSRSGAGSSFSRT   RAMVVKKIKTROCKT   LSYGLSSSSRSGAGSSSFSRT   RAMVVKKIKTROCKT   LSYGLSSSSRSGAGSSSFSRT   LSYGLSSSSRSGAGSSSFSRT   LSYGLSSSSRSGAGSSSFSRT   LSYGLSSSSRSGAGSSSFSRT   LSYGLSSSRSGAGSSSFSRT   LSYGLSSSRSGAGSSSFSRT   LSYGLSSSRSGAGSSSFSRT   LSYGLSSSRSGAGSSSFSRT   LSYGLSSSRSGAGSSSFST   LSYGLSSSRSGAGSSSSRT   LSYGLSSSRSGAGSSSRST   LSYGLSSSRSGAGSSSSRST   LSYGLSSSRSGAGSSSRST   LSYGLSSSRSGAGSSSRST   LSYGLSSSRSG	t		1		codon for peptide		deletion, \-possible nucleotide insertion)
### PASTMSIRGTQKSYKQSTSK RAFKQAAS*QRSGPGFISSS SRVGSSNFRGGLGGGGGGGG MGGITAYTVNQ)SRALLSPL EVDPNIQAYROPEKEQIKTJ KFASFIDEVGFLEQONIML KWSLLDQKTAYRSNMIDN ESYINNLRRQLETLAQELKI, AELGNMQGLVEDFKNNYED NKSTETENEFYLIKKDVDEA NEXTETENEFYLIKKDVDAA NEXTETENEFYLIKKDVAA NEXTETENEFYLIKKDVAA NEXTETENEFYLIKKOM NEXTETENEFYLIKKOM NEXTETENEFYLIKKO					sequence		
### PASTMSIRGTQKSYKQSTSK RAFKQAAS*QRSGPGFISSS SRVGSSNFRGGLGGGGGGGG MGGITAYTVNQ)SRALLSPL EVDPNIQAYROPEKEQIKTJ KFASFIDEVGFLEQONIML KWSLLDQKTAYRSNMIDN ESYINNLRRQLETLAQELKI, AELGNMQGLVEDFKNNYED NKSTETENEFYLIKKDVDEA NEXTETENEFYLIKKDVDAA NEXTETENEFYLIKKDVAA NEXTETENEFYLIKKDVAA NEXTETENEFYLIKKOM NEXTETENEFYLIKKOM NEXTETENEFYLIKKO	23025	53393	IΔ	23159	81	1339	EL SSPRSSESAPSRISAWEGPPAS
RAFKQAAS*QRSGPGTRISSS   SRVGSSNFRGGLGGGGGGGAGA   MGGITA*TVYNQ\SRALLSPL   EVDPNIQA*VRTQEKEQIKTLI   KFASFIPDE*VGFLEQQ\KML   KFASFIPDE*VGFLEQQ\KML   KFASFIPDE*VGFLEQQ\KML   KFASFIPDE*VGFLEQQ\KML   KFASFIPDE*VGFLEQQ\KML   KFASFIPDE*VGFLEQQ\KML   KFASFIPDE*VGFLEQQ\KML   KFASFIPDE*VGFLEQQ\KML   KFASFIPDE*VGFLEQQ\KML   KFASFIPDE*VGFLEQQ\KML   KFASFIPDE*VGFLEQQ\KML   KFASFIPDE*VGFLEQQ\KML   KFASFIPDE*VGFLEQQ\KML   LEALQQTKKLLEMENMSII   KTTSQYAGGGYSLAYGGLTGS   KFASFIPDE*VGFLEQQ\KML   LEALQQTKKLLEMENMSII   KTTSQYAGGGYSLAYGGLTGS   LSYGLSSSSRSGAGSSSSSTSKT.   RAMVVKKIKTRGKLLTLSI   LSYGLSSSSRSGAGSSSSSSTSKT.   RAMVVKKIKTRGKLLTLSI   LSYGLSSSSRSGAGSSSSSTSKT.   RAMVVKKIKTRGKLLTLSI   TYMYYMFLYDSTHGKFHGTY   AENGTL.VINGNITIFIQEQO   KIKWAPILAKVIHDNFGILEGI   TTYMTITATQTINGPSGKLSK   CRALQDIPYSTGAAKPYGK*   ELNGKLTGMAFHVPTANVS   DLTCRLEFPAKYDDIKKNTN   STFDAGAAIVLKDIISVKLISV   DNEFGYSRRVVHLMAHNAS   DNEFGYSRRVVHLMAHNAS   DNEFGYSRRVVHLMAHNAS   DNEFGYSRRVVHLMAHNAS   SAHASAVRQSAAIFFCVASI   HLAETPLGKVKVGVNGFGGIL   LVTRAAFNSGKVDIVANDPI   LNYMYYMFQYDSTHGKFHC   VKAENGKLVINGNPITTIFQQE   PSIKIKWODAGAE*VRRGSFLCFRQFWKAENGKLVINGNPITTIFQQE   PSIKIKWODAGAE*VRRGSFLCFRQFWKAENGKLVINGNPITTIFQQE   PSIKIKWODAGAE*VRRGSFLCFRQFWKAENGKLVINGNPISTTFDAGAGGLKGGSQKG   SAPSAECPPCFVMGCEPMK*   DKQAFKNISNASCTNQLL**H   AKG*FHDNFGNSLEGHD**H   AKG*FH	12023	33333	1	23137	ļ°.	1557	
SRVGSSNFRGGLGGGYGGA   MMGGITAVTVNQNRALLSPL   EVDPNIQAVTVNQNRALLSPL   EVDPNIQAVTVNQNRALLSPL   EVDPNIQAVTVNQNRALLSPL   EVDPNIQAVTVNQNRALLSPL   EVDPNIQAVTQPERQIKTLI   KFASIFIDEVGFLEQQNKML   KWSLLDQQKTARSNMDINN   ESYINNLRRQLETLAQEKLK: AELGNMQGLVEDFKNNYED   NKSTETENEFVLIKKDVDEA   NEXTETENEFVLIKKDVDEA   NEXTETENEFVLIKKDVDEA   NEXTETENEFVLIKKDVDEA   NEXTETENEFVLIKKDVDEA   NEXTETENEFVLIKKDVDEA   NEXTETENEFVLIKKDVDEA   NEXTETENEFVLIKKDVDEA   NEXTETENEFVLIKKDVDEA   NEXTETENEFVLIKKDVEA   NEXTETENEFVLIKKDVEA   NEXTETENEFVLIKKDVEA   NEXTETENEFVLIKKDVEA   NEXTETENEFVLIKKDVEA   NEXTETENEFVLIKKDVEA   NEXTETENEFVLIKKDVEA   NEXTETENEFVLIKKDVEA   NEXTETENEFVLIKKDVEA   NEXTETENEFVLIKKDVEA   NEXTETENEFVLIKKDVEA   NEXTETENEFVLIKKDA   NE	i						
MGGITATTVNQISRALLSPL     EVDPNIQAVRTQEKEQIKTLI     KFASIFIDPEVGFLEQQNKML     KFASIFIDPEVGFLEQQNKML     KFASIFIDPEVGFLEQQNKML     KWISLLDQQKTAVRSNMDINI     ESYINNLRRQLETLAQEKLK     AELGNMQGLVEDFKNNYED     NKSTETENEFVLIKKDVDEA     NELYEEIRELGQGISDTSVV     DNSRSLDTDSIJAEVKTEISEN     RNISWLQAETEGLAGGRISL     GIADAEQRRELVIKDANAKL     LEAALQQTKKLLEMENMSII     KTTSGVAGGVSLAVGGLTG     LSYGLSSSSRSGAGSSSFSRT     RAMVVKKIKTROCKLLTLS     LSYGLSSSSRSGAGSSSFSRT     RAMVVKKIKTROCKLLTLS     LSYGLSSSSRSGAGSSSFSRT     RAMVVKKIKTROCKLLTLS     LSYGLSSSSRSGAGSSSFSRT     RAMVVKKIKTROCKLTLS     LEAALQQTKKLLEMENMSII     KTTSGVAGGVSLAVGGLTG     LSYGLSSSRSGAGSSSFSRT     RAMVVKKIKTROCKLTLS     RAMVVKKIKTROCKLTLS     LSYGLSSSSRSGAGSSSFSRT     RAMVVKKIKTROCKLTLS     CALQDIFVSTHOKHGOKLT     AENGLYJNANDFIDIPOPO     KIKWAPLAKVIHDNFGIEG     TTVFHTLATDJROFSGKLS     CRALQDIFVSTGAAKPVGK     ELNGKLTGMAFHVPTANNS     DLTCRLERPAKVDIKKNIN     DLTCRLERPAKVDIKKNIN     DLTCRLERPAKVDIKKNIN     DLTCRLERPAKVDIK     CALQDIFVSTGAAKPVGK     ELNGKLTGMAFHVPTANNS     DLTCRLERPAKVDIK     DLTCRLERPAKVDIK     VKAENGKLVINONPITTVQE     PSIKIKWQDAGAEVRRGSFL     FNQPWRKAGGSICKGGSKG     SAPSAECPPCFVMGCEPMK     DKAGRALKONIPASTGAAK     GKVIPELNGKLTGMAFRVPT     VSVVDLTCRLERFAKVDIK     VVKQASEGPLKGILGVTEHQ     VSSDFNSDTHSSTFDAGAIG     NDHFFKLISWYDNEFGYSNI     VDLMAHMGFQGS     23029   33397     A   23161   931   1124     1124	1						
EVDPNIQAVRTQEKEQIKTLI   KFASIFIDEVGFLEQONKML   KFASIFIDEVGFLEQONKML   KFASIFIDEVGFLEQONKML   KFASIFIDEVGFLEQONKML   KFASIFIDEVGFLEQONKML   KFASIFIDEVGFLEQONKML   KFASIFIDEVGFPENNYED   ALEIOMOGLIVEPENNYED   ANELYEEJRELQQISDTSVV.   DNSRSLDTDSIIAEVKTEISEN   RNISWLQAETEGLKGGRISLI   GJADAEORRE, VIRDANAKL   LEAALQQTKKLLEMERMSII   KTTSGYAGGVSLAYGGLTG: LSYGLSSSRSGAGSSSFSTR.   RAMVVKIKTRDGKLLTLS.   LYGLSSSRSGAGSSSFSTR.   RAMVVKIKTRDGKLLTLS.   TAMTAFISGKYDIVAINDPRILIEGI   TTYHTITATQTINGPSGKLSR   CRALQDIPVSTAAKPVGK.   RAMVVKIKTRDGKLLTLS.   RAMVVKIKTRDGKLLTLS.   RAMVVKIKTRDGKLLTS.   RAMVVKIKTRDGKLLTLS.   THY THY TATAFISGKYDAMPATAN   STEDAGAAIVTKDISVAINDPRILIEGI   TTYHTITATQTINGPSGKLSR   CRALQDIPVSTGAAKPVGK.   ELNGKLTGMAFHVPTANVS   DLTCRLERPAKYDDIKKNTSTFDAGAAIVKLISV   DNEPGYSNRVVHLMAHNAS   DNEPGYSNRVVHLMAHNAS   DNEPGYSNRVVHLMAHNAS   TSTDAGAAIVKDISVKLISV   DNEPGYSNRVVHLMAHNAS   TSTDAGAAIVKDISVKLISV   DNEPGYSNRVVHLMAHNAS   TSTDAGAAIVKDISVKLISV   DNEPGYSNRVVHLMAHNAS   TSTDAGAAIVKDISVKLISV   DNEPGYSNRVVHLMAHNAS   TSTDAGAAIVKDISVKLISV   DNEPGYSNRVVHLMAHNAS   TSTDAGAAIVKDISVKLISV   DNEPGYSNRVVHLMAHNAS   TSTDAGAAIVKDISVKLISV   DNEPGYSNRVVHLMAHNAS   TSTDAGAAIVKDISVKLISV   DNEPGYSNRVVHLMAHNAS   TSTDAGAAIVKDISVKLISV   TSTDAGAAIVKDISVKLISV   DNEPGYSNRVVHLMAHNAS   TSTDAGAAIVKDISVKLISV   TSTDAGAAIVKDISVKLISV   TSTDAGAAIVKDISVKLISV   TSTDAGAAIVKDNITTTOGE   PSIKKWODAGAEVRRGSPLEFIQPSRRKG	ĺ						
	ĺ						
	l		1				
ESYINNLRRQLETLAGELIX:   AELGNMQGLVEDFKNNYED   NKSTETENEFYLIKKDVDEA   NKSTETENEFYLIKKDVDEA   NKSTETENEFYLIKKDVDEA   NKSTETENEFYLIKKDVDEA   NESTETENEFYLIKKDVDEA   NESTETENEFYLIKKDVDEA   NESTETENEFYLIKKDVDEA   NESTETENEFYLIKKDVDEA   NESTETENEFYLIKKDVDEA   NESTETENEFYLIKKDVDEA   NESTETENEFYLIKKDVBLAKLL   LEALQQTYKKLLEMENMSII   KTTSQYAGGVSLAYGGLTG: LSYGLSSSSRSGASSSSFSTE   RAMVVKIKTRDGKLLTLSI. LEALQQTYKKLLEMENMSII   KTTSQYAGGSUSLAYGGLTG: LSYGLSSSSRSGAGSSSFSTE   RAMVVKIKTRDGKLLTLSI. LEALQQTYKKILETDEA   NESTETENEFYLIKKDY   NES	l		1		· ·	İ	
AELGNMQGLVEDFKNYED	l		1			ŀ	
NKSTETENEFVLIKKDVDEA   NELYEEBIRELQSQISDTSVV   DNSRSLDTDSIIAEVKTEISER   RNISWLQAETEGLKGGRISLI   GJADAGORRE   VIGNANKL   LEAALQOTKKLLEMENMSI   KTTSGYAGGVSLAYGGLTG: LSYGLSSSRSGAGSSSFSRT   RAMVVKIKTRDGKLLTLS.   LEAALQOTKKLLEMENMSI   KTTSGYAGGVSLAYGGLTG: LSYGLSSSRSGAGSSSFSRT   RAMVVKIKTRDGKLLTLS.   RAMVVKIKTRDGKLLTLS.   MTAFNSGKVDIVAINDPGILGG   TTVHTITATQTINGPSGKLSR   CRALQDIPVSTAAKPYGK*   RENGILVINGNPITIFQDQD   KIKWAPILAKVIHDNFGILEG   TTVHTITATQTINGPSGKLSR   CRALQDIPVSTAAKPYGK*   ELNGKLTGMAFHVPTANVS   DLTCRLEFPAKYDDIKKNT   STFDAGAAIVLKDHSVKLISV   DNEFGYSNRVVHLMAHNAS   DNEFGYSNRVVHLMAHNAS   DNEFGYSNRVVHLMAHNAS   LYTRAAFNSGKVDIVAINDPI   LNYMYYMFQVDSTHGKFHC   VKAENGKLVINONPITTROE   PSIKIKWODAGAEYRRGSFL   FNOPWRKAGGSICKGGSGK   ENGRAF   FNOPWRKAGGSICKGGSKK   ENGRAF   ENGRA	1		1				
NELVEEEIRELQSQISDTSVV			1				
DNSRSLDTDSILAEVKTEISER   RNISWLQAETEGLKGGRISLI   GIADAEQRREL.VIKDANAKL   LEAALQQTKKLLEMEMMSII   KTTSGYAGGGVSLAYGGLTGS   LSYGLSSSSRSGAGSSSSSTSRT   RAMVVKKIKTROKLLTLSL   LSYGLSSSSRSGAGSSSSSTSRT   RAMVVKKIKTROKLLTLSL   LSYGLSSSSRSGAGSSSSSTSRT   RAMVVKKIKTROKLLTLSL   LSYGLSSSSRSGAGSSSSTSRT   RAMVVKKIKTROKLLTLSL   LSYGLSSSSRSGAGSSSSTSRT   RAMVVKKIKTROKLLTLSL   LSYGLSSSSRSGAGSSSSSTSRT   RAMVVKKIKTROKLTLSL   LSYGLSSSSRSGAGSSSSSTSRT   RAMVVKKIKTROKLTLSL   LSYGLSSSSSSSTST   RAMVVKKIKTROKLTLSL   LSYGLSSSSSSSSTST   RAMVVKKIKTROKLTLSL   LSYGLSSSSSSSSTST   RAMVVKKIKTROKLTLSL   LSYGLSSSSSSSST   LSYGLSSSSSSSST   LSYGLSSSSSSSST   LSYGLSSSSSSSST   LSYGLSSSSSSSSST   LSYGLSSSSSSSSST   LSYGLSSSSSSSSST   LSYGLSSSSSSSSSSST   LSYGLSSSSSSSSSST   LSYGLSSSSSSSSSSST   LSYGLSSSSSSSST   LSYGLSSSSSSSSSST   LSYGLSSSSSSSSSSST   LSYGLSSSSSSSSSSST   LSYGLSSSSSSSSSSSSSSSSSSST   LSYGLSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSS						ŀ	
RNISWLQAETEGLKGGRISLI   GIADAEQRRELVIKDANAKL   LEAALQQTKKLLEMENMSI    KTTSGYAGGVSLAYGGLTG    LSYGLSSSSRSGAGSSSFSTR RAMVVKIKTRDGKLLTLSI   LSYGLSSSSRSGAGSSFSTR RAMVVKIKTRDGKLLTLSI   LSYGLSSSRSGAVSFSTR RAMVVKIKTRDGKLLTLSI   MTAFNSGKVDIVAINDFEIDE    THAT PART SCHOLLT SI   AENGLIT JINGPISGKLSR     CRALQDIPVSTGAAKPVGST     ELNGKLTGMAFHVPTANVSI     DLTCRLERPAKYDDIKKNIT     STFDAGAAIVLADISVALISI     DNEFGYSNRVVHLMAHNAS     DNEFGYSNRVVHLMAHNAS     A 23162 3 1151     ASAHASAVRGSAIFFCVASI     HIAETPLGKVKVGVNGFGRI     LVTRAAFNSGKVDIVAINDPI     LNYMVYMFQVDSTHGKFHC     VKAENGKLVINONIPTITOCE     PSIKIKWOIDAGAEVRRGSPL     FNQPWRKAGGSICKGGSQKG     ISAPSAECPPC-VMGCEPMKH     DKQAFKNISNASCTNQLL-H     AKGPHIDNFGNSLEGLIDH     MATTANPRKTVDIGPSRGKL     LMGRGALJQNIIPASTGAAK, GKVPELNGKLTGMAFRVPT     VSVVDLTCRLERPAKYDDIK     VVKQASEGPLKGLIGVTERG     VSDFNSDTHSSTFDAGAIG,     NDHFVKLISWVJDNEFGYSNI     VVDLMAHMGFQGS     23029   33397     A 23163   160   236     2360     SSJPSS     A 23164   931   1124							
GJADAGORREL.VIKDANAKL   LEAALQQTKKLLEMENNSII   KTTSQYAGGVSLAYGGLTG   LSYGLSSSSRSGQSSSSFSRT   RAMVVKIKTROCKLLTLS   LSYGLSSSSSSFSRT   RAMVVKIKTROCKLLTLS   LSYGLSSSSRSGAGSSSFSRT   RAMVVKIKTROCKLLTLS   LSYGLSSSSRSGAGSSSFSRT   RAMVVKIKTROCKLTLS   LSYGLSSSRSGAGSSSFSRT   RAMVVKIKTROCKLTLS   LSYGLSSSRSGAGSSSFSRT   RAMVVKIKTROCKLTLS   LSYGLSSSRSGAGSSSFSRT   RAMVVKIKTROCKLTLS   LSYGLST   LSY	ĺ						
LEAALQOTKKLLEMENMSII   KTTSGYAGGVSLAYGGLTG: LSYGLSSSSRSGAGSSSFSRT   RAMVVKIKTTDGKLLTLS!   LSYGLSSSSRSGAGSSSFSRT   RAMVVKIKTTDGKLLTLS!   RAMVVKIKTTDGKLLTLS!   MTAFNSGKVDIVAINDPFID   YMYWHFLYDSTHGKFHGTY   AENGTLVINGNPITIFQDQDG   KIKWAPILAKVHIDNFGHIEGI   TITVHTITATQTINGPSGKLSR   CRALQDIIPYSTGAAKPYGK*   ELNGKLTGMAFHVPTANVS!   DLTCRLEKPAKVDDIKKINT   STFDAGAAIVLKDHSVKLISV   DNEFGYSNRVVHLMAHNAS   DNEFGYSNRVVHLMAHNAS   DNEFGYSNRVVHLMAHNAS   TSTPAGAAIVLKDHSVKLISV   DNEFGYSNRVVHLMAHNAS   LVTRAAFNSGKVDIVAINDPI   LNYMYYMFQVDSTHGKFHG   VKAENGKLVINONPITTROE   PSIKIKWODAGAEYRRGSFL   FNOPWRKAGGSICKGGSGKG   ESAPSAECPPCFVMGCEPMKI   DKQAFKNISNASCTNQLL*H   AKG*FHDNFGNSEGLIDH*   MAITANPRKTVDIGPSRGKL   LMGRGALCONIPASTGHAK   GKVIPELNGKLTGMAFRVPT   VSVVDLTCRLEKPAKYDDIK   VVKQASEGPLKGLIGAYTERQ   VSDFNSDTHSSTFDAGAIG, NDHFVKLISWYDDNEFGYSNI   VVKQASEGPLKGLIGAYTERQ   VSSDFNSDTHSSTFDAGAIG, NDHFVKLISWYDDNEFGYSNI   VDLMAHMGFQGS   23029   33397   A   23163   160   236   236   2360   2					ł		
23026   53394   A   23160   1   651   MTAFNSGKYDIVAINDPFIDL   YMYMFLYSTHERMEN   A   23160   1   651   MTAFNSGKYDIVAINDPFIDL   YMYMFLYSTHERMEN   A   23160   1   651   MTAFNSGKYDIVAINDPFIDL   YMYMFLYSTHERMEN   A   A   A   A   A   A   A   A   A			1			ļ	
LSYGLSSSRSGAGSSSPSRT.   RAMVVKKIKTROKLLTLSI			1				
RAMVVKIKITROCKLITLSI   23026   53394   A   23160   1   651   MTAFNSGKYDIVAINDPFIDI   YMVYMFLYDSTHGKFIGTY   AENGTLVINGNPITHFQDQDI   KIKWAPLAKVHIDNGBIGEI   TTVHTITATQTINGPSGKLSR   CRALQDIPVSTGAAKPVGK   ELNGKLTGMAFHVFTANVSI   DLTCRLEKPAKVDDIKKNIT   STFDAGAAIVKDBISVKLISI   DNEFGYSNRVVHLMAHNAS   23028   53396   A   23162   3   1151   ASAHASAVRQSAAIFFCVASI   HAETPLGK VKVGVNGFGRI   LVTRAAFNSGKVDIVAINDPI   LNYMVYMFQVDSTHGKFHC   VKAENGKLVINGNPITHFOE    PROPWRKAGGSICKGGSQKC   ISAPSAECPPCFVMGCEPMKI   DKQAFKNISNASCTNQLL*H   AKG*FHDINFGNSLEGILDHI   MAITANPRKTVDIGPSRGKL   LMGRGALQNIIPASTGVAAK   GKVPELNOKLTGMAFRVFT   VSVVDLTCRLEKPAKVDDIK   VVKQASEGPLAGGIGVCGIGVAFT   VSVVDLTCRLEKPAKVDDIK   VVKQASEGPLAGGIGVTERICATION   VKKANINAMATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPLAGGIGVTERICATION   VVKQASEGPL			1	i			
23026   53394   A   23160   1   651   MTAFNSGKYDIVAINDFFIDI			1		ì		
YMYYMFLYDSTHGKFHGTY	22026	52204	<u>                                     </u>	22160	,	651	
AENGTLVINGNPITHEQDOM   KIKWAPILAKVIHDNFGIIEGI   TTVHTITATQTINGPSGKLSR   CRALQDIIPVSTGAARPVGK*   ELNGKLTGMAFHYPTANNS*   DLTCRLEKPAKYDDIKKINT STEDAGAAIVLKDIISVKLISV   DNEFGYSNRVVHLMAHNAS   DNEFGYSNRVVHLMAHNAS   DNEFGYSNRVVHLMAHNAS   DNEFGYSNRVVHLMAHNAS   S3396   A 23162   3   1151   ASAHASAVRQSAAIFFCVAS   HIAETPLGKVEVGVNGFGRI LVTRAAFNSGKVDIVAINDPI LNYMYVMFQVDSTHGKFHE VKAENGKLVINGNPITHFGE   PSKIKWGDAGAEYRRGSPLGFNQFWRKAGGSICKGGSQKG   ISAPSAECPPCFVMGCFPMKF DKQAFKINSNASCTNQLL+H AKG*PHDNFGNSLEGLHDHK MAITANPRKTVDGPSRGKLV   LMGRGALKONIPASTGAAK, GKVPELNGKLTGMAFRVPT VSVVDLTCRLEKPAKYDDIK VVKQASEGPLKGILGYTEHQ VSSDFNSDTHSSTFDAGAIG, NDHFVKLISWYDDNEFGYSNI VVDLMAHMGFQGS   23029   33397   A 23163   160   236	23020	33374	<u> </u> ^	23100	ľ	051	
	ļ	1	1				
TTYHTITATQTINGPSGKLSR   CRALQDIIPVSTGAAKPVGK   ELNGKLTGMAFHVPTANVS   DLTCRLEKPAKYDDIKKNT   STFDAGAAIVKDIISVKLISV   DNEFGYSNRVVHLMAHNAS   DNEFGYSNRVVHLMAHNAS   33395   B   23161   1   382     ASAHASAVRQSAAIFFCVASI   HIAETPLGK VEVGVNGFGRI   LVTRAAFNSGKVDIVAINDPI   LNYMYMFQVDSTHGKFHC   VKAENGKLVINONPITIFOED   PSIKIWOIDAGAEVRRGSPL   FNQPWRKAGGSICKGGSQKC   ISAPSAECPPCFVMGCEPMKH   DKQAFKNISNASCTNQLL*H   AKG*FHDINFGNSEGLIDH   MAITANPRKTVDIGPSRGKL   LMGRGALQNIIPASTGVAAK   GKVIPELNOKLTGMAFRVPT   VSVVDLTCRLEKPAKYDDIK   VVQASEGPLKGLIGVTERIQ   VSDFNSDTHSSTFDAGAIG   NDHFVKLISWYDNEFGYSNI   VVQASEGPLKGLIGVTERIQ   VSDSPNSDTHSSTFDAGAIG   NDHFVKLISWYDNEFGYSNI   VDLM\()AHMGFQGS   23029   33397   A   23163   160   236     236     2360     2361							
CRALQDIIPVSTGAARPVGK:   ELNGKLTGMAFHVPTANVS'     DLTCRLERPARYDDIKKNT'     STPDAGAAIVLKDIISVKLISV     DNEFGYSNRVVHLMAHNAS     DNEFGYSNRVVHLMAHNAS     S3396			1				
ELNÖKLTGMAFHVPTANVS							
DLTCRLERPAKYDDIKKINT			l				
STFDAGAAIVLKDIISVKLISV			1		l		
DNEFGYSNRVVHLMAHNAS			1				
23027   53395   B   23161   I   382			1				
23028   53396   A   23162   3   1151	23027	53395	В	23161	1	382	Did of Grace Characteristics
LVTRAAFNSGKVDIVAINDP    LNYMYYMFQYDSTHGKFHC   VKAENGKLVINONPITTROEL    PSIKIKWOIDAGAEVRRGSFLE    FINOPWRKAGGSICKGGSQK   ISAPSAECPPCFVMGCEPMK    DKQAFKIISNASCTINQLL*H   AKG*FHDNFGNSLEGLIDHIK   MAITANPRKTVDIOPSRGKLI   LMGRGALLOIMPBASTGAAK,   GKVIPELNGKLTGMAFRVPT   VSVVDLTCRLEKPAKYDDIK   VVKQASEGPLKGLIGVTTERIQ   VSDFNSDTHSSTFDAGAIG,   NDHFVKLISWYDDNEFGYSNI   VDLM\  AHMGFQGS   23029   33397   A   23163   160   236     23029   33398   A   23164   931   1124			A		3	1151	ASAHASAVROSAAIFFCVASRA
L.NYMYYMFQYDSTHGKFHC							HIAETPLGKVKVGVNGFGRIGR
L.NYMYYMFQYDSTHGKFHC			1				LVTRAAFNSGKVDIVAINDPFID
VKAENGKLVINONPITIFOE   PSKIKWGDAGAEYRRGSPLG    FNQPWRKAGGSICKGGSQKG    FNQPWRKAGGSICKGGSQKG    ISAPSAECPPCFVMGCFPMK    DKQAFKINISMASCTINQLI-H    AKG*FHDNFGNSLEGLHDHK    MAITANPRKTVDGPSRGKLV    LMGRGALKONIPASTGAKA,   GKVIPELNGKLTGMAFRVPT    VSVVDLTCRLEKPAKYDDIK    VVKQASEGPLKGILGYTEHQ    VSSDPNSDTHSSTFDAGAIGI,   NDHFVKLISWYDDNEFGYSNI    VDLM\(\text{AHMGFQGS}\)   23029   33397   A   23163   160   236     23030   33398   A   23164   931   1124			1				LNYMVYMFQYDSTHGKFHGT
FNQPWRKAGGSICKGGSQKC   ISAPSAECPPCFVMGCEPMKI   DKQAFKNISNASCTNQLL+H   AKG*FHDNFGNSLEGLHDHK   MAITANPRKTVD/GPSRGKL\							VKAENGKLVINGNPITI\FQE\RD
FNQPWRKAGGSICKGGSQKC   ISAPSAECPPCFVMGCEPMKI   DKQAFKNISNASCTNQLL+H   AKG*FHDNFGNSLEGLHDHK   MAITANPRKTVD/GPSRGKL\	i				l		PSKIKWG\DAGAEYRRGSPLGV
ISAPSACPPCFVMGCEPMK#   DKQAFKNISNASCTNQLL*H   AKG*FHDNFGNSLEGLHDH#   MAITANPRKTVDIOPSRGKL    LMGRGALKOJIPBASTGVAK,   GKVIPELNGKLTGMAFRVPT     VSVVDLTCRLERPAKYDDIK     VVKQASEGPLKGLIGVTTERIJE     VSDFNSDTHSSTFDAGAIG,     NDHFVKLISWYDMEFGYSNI     VJLMVAHMGFQGS     23029   33397   A   23163   160   236     23030   53398   A   23164   931   1124	1		1				FNOPWRKAGGSICKGGSQKGSS
AKĞPHDNEGNSLEGLİIDIK   MAITANPRKTVDIĞPSRĞKLÜ   LMĞRĞALÜNIPASTĞYAK, GKVIPELNÖKLTĞMAFREVPT VSVVDLTCKLEKPAKYDDIK VVKQASEĞPLKĞILĞYTEHQ VSSDFNSDTHSSTFDAĞAĞI, NDHFVKLISWYDNEFĞYSNI VDLM'AHMĞFQĞS	1		1				ISAPSAECPPCFVMGCEPMKKY
AKĞPHDNEGNSLEGLİIDIK   MAITANPRKTVDIĞPSRĞKLÜ   LMĞRĞALÜNIPASTĞYAK, GKVIPELNÖKLTĞMAFREVPT VSVVDLTCKLEKPAKYDDIK VVKQASEĞPLKĞILĞYTEHQ VSSDFNSDTHSSTFDAĞAĞI, NDHFVKLISWYDNEFĞYSNI VDLM'AHMĞFQĞS	1		1				DKQAFKI\ISN\ASCTNQLL*HPL
MAJTANPRKTVD/GPSRGKL  LMGGAL/QNIIPASTG/AAK, GKVIPELNGKLTGMAFRVPT VSVVDLTCRLEKPAKYDDIK VVKQASEGPLKGILGYTEHQ VSSDFNSDTHSSTFDAGAIG, NDHFVKLISWYUDNEFGYSNI VIDLM/AHMGFQGS   23029   33397   A   23163   160   236     2360   33398   A   23164   931   1124			1				
LMGRGAL\QNIIPASTG\AAK.   GKVIPELNGKLTGMAFRVPT   VSVVDLTCRLERPAKYDDIK   VVKQASEGPLKGILGYTEHQ   VSSDTSDTHSSTFDAGA\GI.   NDHFVKLISWYDNEFGYSNI   VDLMAHMGFQGS   23029   53397   A   23163   160   236   236   2303   33398   A   23164   931   1124			1				MAITANPRKTVD\GPSRGKLWP
GKVIPELNGKLTGMARRVPT   VSVVDLTCRLEKPAKYDDIK   VVKQASEGPLKGILGYTEHQ    VSSDPNSDTHSSTFDAGAGI,   NDHFVKLISWYDDMEGYSNI   VDLM\AHMGPQGS   23029   33397   A   23163   160   236   236   23030   33398   A   23164   931   1124			1				LMGRGAL\QNIIPASTG\AAKAV
VSVVDLTCRLEKPAKYDDIK VVKQASEGPLKGILGYTEHQ VSSDFNSDTHASTFDAGA/GI, NDHFVKLISWY\DNEFGYSNI V\DLM\AHMGFQGS   23029   53397 A   23163   160   236     23030   53398 A   23164   931   1124	1	1	1		l		GKVIPELNGKLTGMAFRVPTAN
VSSDFNSDTHSSTFDAGA\GI, NDHFVKLISWYIDNEFGYSNI V\DLM\AHMGFQGS   23029   53397   A   23163   160   236     23030   53398   A   23164   931   1124					ŀ		VSVVDLTCRLEKPAKYDDIKK
VSSDFNSDTHSSTFDAGA\GI, NDHFVKLISWYIDNEFGYSNI V\DLM\AHMGFQGS  23029 53397 A 23163 160 236 23030 53398 A 23164 931 1124	1		l				
NDHFVKLISWY\DNEFGYSNI V\DLM\AHMGFQGS   23029   53397   A   23163   160   236       23030   33398   A   23164   931   1124		1	1		1		VSSDFNSDTHSSTFDAGA\GIAL\
VIDLM\AHMGFQGS 23029 53397 A 23163 160 236 23030 53398 A 23164 931 1124	1		1		1		NDHFVKLISWY\DNEFGYSNRV
23029 53397 A 23163 160 236 23030 53398 A 23164 931 1124							
23030 53398 A 23164 931 1124	23029	53397	Α	23163	160	236	
בטכבון בטכבון בטכבון בטכבון בטכבון בטכבון בטכבון	23031	53399	Α	23165	132	303	

	ptide ho		location of first	codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
		09/540,217	codon for peptide	of peptide sequence	deletion, \"possible nucleotide insertion)
			sequence		
23032 534	00 4	23166	120	1171	
		23167	1008	1262	HELOCSLLPVRHATAGHTNGQ
23033 534	01  A	23107	1008	1202	NRLHWGHVAHSNYGAALKTN
			1		GVEVR*FTAAIARCDGNPYESF
		20160	700	11/0	RKDVRADFQTRIDAIVRCLPK
23034 534	02 A	23168	732	1162	RCHEKGRKGHQAPTGSTRPDG
					TLSACRSCQQGQGMPASAGVP
- 1			1		ARQMGSTGNLPHLV*PYFLGLS
				<u> </u>	SKAQHTG*LHKKR
23035 534		23169	2	1093	
23036 534		23170	236	3084	
23037 534					LRDGRGSMSSSSPNSASGSAICP
23038 534	06 A	23172	305	522	
	i				ASSVMPETVSVSRVMPNVTKP
					A/MTAKQTRIRNQMVAITYIES
				***	DAWNRSYQ
23039 534		23173	177	386	CANADIANNI ANDVOCA A A CODO
23040 53408	08 A	23174	3	200	GAVAPMHPLAHPVPGAAAQPP PA*QPATPCHQPA*GQPAPPDV
		20105	<del>   </del>	101	PGEPSAGRSLHHAHICLAPNEQ
23041 534		23175	171	535	LIPTORIOTEL TOLICUPE OFFICE
23042 53410	10 A	23176	190	505	VPTDRIRTRLTSVCKRRCEVHQ
	- 1				LEDRPFEVYQVA/PRRD*SFETS
	- 1		1		VWNHMRKQ/PPLDVRQLAS*N
		1			VLHRQLDIMDSAVASMYSIHLS
			0.10	1070	IKMGIGFSGVTTVVKGTPK
23043 534		23177	349 518	1272	VSSSIIMFGPLSQFPRQQVGSSV
23044 534	12 A	23178	218	1451	KRPSNVVCPGLMFSFFSSAA*IP
		1			**PLI*OPGV
					TSPDVLRAGPRVPSVOGTSPSO
23045 534	13 A	23179	2436	2894	
	- 1				SCPGQRPHGPNSKAYGR/LSVM
	- 1			ŀ	AQYYCNVIPVLE/VPPSAFTPPP
					KVDSAVV/RLVPHATMPHPVK
				i .	DVRVLSRITTEAFNQRRKTIRNS
				į .	LGNLFSVEVLTGMGIDPAMRA
					ENISVAQYCQMANYLAENAPL
23046 534	14 A	23180	1846	2557	IPVSERQGSDWRFFSVYVVNKA
					GGLIYQLDSYAPTGLRLEKTF\S
		1			YPLDLLLKLHDERVLVAFGQR
				1	GRHPEWGHAVL\AINGMDVNG
					RYTAD\GKKVLEYLG\NP\ANYP
			ı		V\SYLFAQPRL\TSNDKLMLAL
			1		MFHSLFAIGSQACLP\EQGKPQG
	1		I		IENAWKTDTFKLH\CYQTLT\GI
		1			KFVVLADPRQAGIDSLLRKIYEI
			1	1	YSDFALKNPFYSLEMPIRCELFD
					QNLKLALEVAEKAGTFGPGS
23047 534		23181	34	225	
	16 B	23182	105	381	l .
23048 534 23049 534		23182	1	837	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ł	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		<u> </u>			744	LIE + GODGERD DOLL + DEDENIO
23050	53418	A	23184	3	/44	HEASCRSERRRQMADFDTYDD
1						RAYSSFGGGRGSRGS\AGGHGS
ľ						RSQKELPTEPPYTAYVGNLPF\N
						TVQGDIDAIFKDLSIRSVR\LVR
						DKVTGKFK\GFCYVEFDEV\DSL
			1			K\EALTYD\GALLGDRSL\RVDF
l						AKGRKTKIRVAFGF\RKGGPDD
	1					R\GFRDDF\LGGRGGSRPGD\RR
						TGPPMGSR\FRD\GPPLRG\SN\M
1			1			DFREPTEEERAQRPRIQ\LKPRT
						VWRPLNQVANPNSAIFGGARPR
ļ						EEVVQKEQE
23051	53419	С	23185	1	1038	
23052	53420	Α	23186	1	1486	MGRNQSRKAENSKHESTYSPP
		1				KDHSSSQAMEQSWTENGFEKL
						GFRRKEALYYLDLTASCQSQEL
	ľ	1				FQLYAGMSVVGTSMPVQAVCP
						YCGNRIITVTTFVPGALTWLLC
		ĺ			İ	TTLFLFGYVLGCCFLAFCIRSLM
	ľ	l				DVKHSCPVCQRELFYYHRLRQ
	ŀ					MELASSARTVVAPSRPCAGREG
	İ				ŀ	RGDANGKNSSPTAGSAMSSEPP
1	ŀ					PPPQPPTHQA\GVGLLDTPRSRE
			l .	l .		RSPSP\LRGNVVP\SPLPTRRTRT
			1			FS\ATVR\ASQGPVYKGV\CKCF
		ļ	l			CRSKG/HIGFNYPPAGWAGPDIF
	İ	l				PGTSLDVGRGSY\VPSGKGDEV
	ŀ	1				TLLKCAPIPTPKNE\KLAGPWEV
		1				VHHSPWHPGT\KHET\WSG\HVI
		1			i	SFLGGLVESTPLSLCFVGDFAG
		1				RROOTLEDDILPHETGLORGNG
		1				PSQVSPGGKGYGGAGVGCGVF
	İ	1			1	PAISTAYGPLOOPLHHLKSIKSH
		1	ŀ			LKRRGAHWWLSGGSNPIPGSGS
		1				RVGISSQLAQTHGLNPQNPSSS
	i	1				
22052	62421	<u> </u>	23187	2	617	WSWRPQIPRSGPLAA
23053	53421	A_		1	993	
23054	53422	A	23188	779	1210	A A A COST A CRIDED CA DISTRICTURA
23055	53423	Α	23189	1/19	1210	AAAGSEAGRRRGADPTRGPHV
				1		LPGSRQRRP/PIFPSLDLSGFFLF
	1			İ		CSEFCPEIKSTNPGISIGDVAKE
	1					KKKLGEMWNSLNDSEKQPYFT
						KAAKLKEKYEKDVADYKSKG
	1					KFDGAKGPAKVARKKVEEEDE
		<u> </u>				QEEEEEEEEEEDE

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nuclcotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23056	53424	A	23190	116	522	KAPYSTNSLSLPPFPILFMGSGY
23030	33424	ľ`	25150	1		RKGPTFQEPKPLPGAAARGPGC
		l				DPTQPPRHSAPRDHGPP\SPRCS
		l				PRPARPRASPAWLPPRAPTPPSP
		l	Į.			PAPAAAPRSCRGHARASSGRSG
		1	l			TTATPRRRRGGLWNQSPMDIK
		ı				G
23057	53425	A	23191	1	152	
23058	53426	A	23192	1	676	MWSAGRGGAAWPVLLGLLLA
22000	33 120	ľ.	25152	l'		LLVPGGGAAKTGAELVTCGSV
						LKLLNTHHRVRLHS\HDI\TYGS
		ı				GSGOO\SVTGVEASDDANSYW
		1				RIRGGSEGGCPRGSPVRCGOAV
Į.	i	l		l .		RLTHVLTGKNLHTHHFPSPLSN
1	l					NOEV\SAFGE\EGEGDDPGTLW
						TVRCSGTA/HWEREAAVRFQH
	İ	1	İ			VGTF\VFLSVTGEQYGSPIRGOH
						EVHGMPQCQ\QHNTWKAMEGI
						FIKPSVEPSAGHDEL
23059	53427	Α	23193	1	1044	
23060	53428	Α	23194	1	1083	MTATLAAAADIATMVSGSSGL
			i			AAARLLSR\SFLLPHNGIRHCAY
		l				TA/SLRQHLYVDKNTKIICQGFT
ł		i				GKQG/TPFHSQQALEYGTKLVG
		1				GTTPGKGGQTHLGLPVFNTVK
		1				EAKEQTGATG\SVIYVPPPFAAA
ĺ			1			AINEAIE\AEIPLVVC\YPRGIPQP
		1				GLGHSSSHKTCGPGKDKAYWG
	l	1	ľ			PNCPGV\INPGECKIGIMPG/HIF
	i				i	HKKGRIGIVSTSGTLTYEAGH/Q
1		l			1	TTQKLGLGQSLCVGIGGDP\FN
						GTDFID\CLEILFERFLPQEGIILI
					[	G\EIGGNA\EENAAEILEPLYSGS
		L				KF/ARPVVS\FIAGLTAPPGRRN
					1	GVMPGAIISGREKVGAKRERSS
l	1		1			ALQSAGVVV\SMSPCTAGEPRF
l		1	1		1	YKEFEKREDRL

NO.   of peptide   sequence   load     on USSN   ocodo for peptide   sequence   of peptide   sequence   of peptide   sequence   of peptide   sequence   of peptide   sequence   of peptide   of pepti	SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
23061   53429   A   23195   1173   2110   FFTELEETILKFIWNQKRACISK		of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ARVNTNOKSGGITLPOFKLYNE		- Interest					
AIVAKITWAWYSSRIHOQEKR   REKK*KKDRSVPETLKKKQR   NFTELNIQCLSKKFAQKMQIA   RRKLIYEKVKIICHKEYRKMYR   TEIRMARMATKAGNTYVPAEP   KLAFVIRIGINGVSPKVRKVLQ   LFLLYQIFINGAFVKLMASINM   LRIVEPYIA/WRYPNLKSVNELI   YKCGYGKINKK*IALTDNALTA   RSLGKYGIICREDLHEITYTOK   CFKEANNFLWPF/KLSSPRGGM   KKKTHIFVEGGDAGKREVQIN   RLIRRMN   RLIRRMN   RIFER   RKKRAFAELKIKELRKKEVPAVPETLK   KKRRNFAELKIKELRKKEVPAVPETLK   KKRRNFAELKIKELRKKEVPAVPETLK   KKRRNFAELKIKELRKKEVPAVPETLK   RKKRNFAELKIKELKKEVPAVPETLK   RKKRNFAELKIKELRKKEVPAVPETLK   RKKRNFAELKIKELKEKKEVPAVPETLK   RKKRNFAELKIKELKEKKEVPAVPETLK   RKKRNFAELKIKELKEKKEVPAVPETLK   RKKRNFAELKIKELKEKKEVPAVPATLK   RKKRNFAELKIKELKKEKKEVPAVPATLK   RKKRNFAELKIKELKKEKKEVPAVPATLK   RKKR	23061	53429	Α	23195	1173	2110	FFTELEETILKFIWNOKRACISK
AIVAKITWAWYSSRIHOQEKR   REKK*KKDRSVPETLKKKQR   NFTELNIQCLSKKFAQKMQIA   RRKLIYEKVKIICHKEYRKMYR   TEIRMARMATKAGNTYVPAEP   KLAFVIRIGINGVSPKVRKVLQ   LFLLYQIFINGAFVKLMASINM   LRIVEPYIA/WRYPNLKSVNELI   YKCGYGKINKK*IALTDNALTA   RSLGKYGIICREDLHEITYTOK   CFKEANNFLWPF/KLSSPRGGM   KKKTHIFVEGGDAGKREVQIN   RLIRRMN   RLIRRMN   RIFER   RKKRAFAELKIKELRKKEVPAVPETLK   KKRRNFAELKIKELRKKEVPAVPETLK   KKRRNFAELKIKELRKKEVPAVPETLK   KKRRNFAELKIKELRKKEVPAVPETLK   RKKRNFAELKIKELKKEVPAVPETLK   RKKRNFAELKIKELRKKEVPAVPETLK   RKKRNFAELKIKELKEKKEVPAVPETLK   RKKRNFAELKIKELKEKKEVPAVPETLK   RKKRNFAELKIKELKEKKEVPAVPETLK   RKKRNFAELKIKELKEKKEVPAVPATLK   RKKRNFAELKIKELKKEKKEVPAVPATLK   RKKRNFAELKIKELKKEKKEVPAVPATLK   RKKR							
REEKKE*KKDPSVPETLK*KOR   NFTELKIQCLSKKFAQKMLQTA   RRKLIVEKVKIICHKEYTKMYR   TEIRMARMATKAQNFYVPAEP   KLAFVIRIGINGVSPKVRKVU, L   LFLLYQIFNGAFVKLNKASINM   LRIVEPYIA/WKYPNLKSVNELI     YKCQYGKINKK*IALTDNALTA   RSLGKYGIICREDLIHEIYTVGK   CFKEANNFLWPF/KLSSPRGGM   KKKTTIIFVEGGDAGKREVQIN   RLIRRINN     23062   53430   A   23196   3   571   GTMEOVEEKKKEVPAVPETLK   KKRRNFAELKIKRI.RKFAQK   MLRKARKLIVEKATA   KKRRNFAELKIKRI.RKFAQK   MLRKARKLIVEKAFAQK   MLRKARKLIVEKAFAQK   MLRKARKLIVEKAFAQK   MLRKARKLIVEKAFAQK   PVVPARTQIWRFVIRINGSINGS   EPQRFRKVLQLLRPFVIRINGSINGS   EPQRFRKVLQLLRPFVIRINGSINGS   GNQFDLHFPTEKRSHPGSQ   QNQFDLHFPTEKRSHPGSQ   QNQFDLHFPTEKRSHPGSQ   QNQFDLHFPTEKRSHPGSQ   QNGFDLHFPTEKRSHPGSQ   QYPVKDKKLLEVCTGGSPSPW   ULMGOSSPSGIFGAFQRGLLR   GTYNKYINVKKGSDLGGFTM   LACYVLFSYSFSYKHLKHER!   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRGGRVLWM   QACIMREYKLVVLGSGGVGKS   ALTVQFVQGIFVEKYPPTIEDS   YRKQVEVDAQQCMLELLDTAG   TEQFTAMRDLYMKNOQGFALV   VSTTAQSTFNIOLQDLREQURV   KDTIDVPMILVGNKCDLEDER   VVGKEQQQNLARQWNNCAFL   ESSAKSKIVNEIFYDLVRQINR   KTYPVPGKAPQKSPSCQLL   ESSAKSKIVNNEIFYDLVRQINR   KTYPVPGKAPQKSPSCQLL   ESSAKSKIVNNEIFYDLVRQINR   KTYPVPGKAPQKSPSCQLL   SPHGEPLLQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCRREMWGWSLH   TESPLCQGCREMWGWSLH   TESPLCQGCREMWGWSLH   TESPLCQGCREMWGWSLH   TESPLCQGCREMWGWSLH   TESPLCQGCREMWGWSLH   TESPLCQGCREMWGWSLH   TESPLCQGCREMWGWSLH   TESPLCQGCREMWGWSLH   TESPLCQGCREMWGWSLH   TESPLCQGCREMWGWSLH   TESPLCQGCREMWGWSLH   TESPL							
NFTELKIQCLSKKFAQKMLQTA   RRKLIYEKVKHCHKEYRKMYR   RRKLIYEKVKHCHKEYRKMYR   TEIRMARMATKAGNIYYPAEP   KLAEVIRIRGINGVSPKVRKVLC   LFLLYQIFINGAFVKLNKASINM   LRIVEPYIAWRYPNLKSVMELI   YKCQYGKINKK*IALTDNALTA   RSLGKYGIICREDLIHEIYTYGK   CFKEANNFLWPF/KLSSPRGGM   KKKTHIFVEGGDAGKREVQIN   RJIRRMN   GTMEGVEKKKEVPAVPETKA   KKRRNFAELKIKRLRKKFAQK   MM.RKARRKLIYEKAKHYHKEY   RQMYRIJERMAJMARKAGN   FIVYPARTQIWREVIRINGRINGS   EPQRFRKVQLQLRPFVLIFNGT   FPVKLQQGFRINMLRDCRSHIL   AWGYPNLKSVHELIYKJRGYG   QNQPLHFPTEKRSHPGSQ   QNQPLHFPTEKRSHPGSQ   QYPVEDKKLLEVGTGGSPFSW   ULMRGPSSPSGIFGVAFQGLR   GTYYNKYINNYKKGSDLGGFTMV   LACVYLFSYSPSYHILKHERE   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRYH						Į.	
RRKLIPEKVKHCHKEYRKMYR TEIRMARMATKAGNFYVPAEP   KLAFVIRIRGINGVSFKVRKVLQ   LFLLYQIFINGAFYKLNKASINM   LRIVEPYIJAWRYPILKSVMELI   YKCGYGKINKK*IALTDNALTA   RSLGKYGIICREDLIHEIYTVGK   CFKEANNFLWPFYLSVSPRGGM   KKKTTIIFVEGGDAGKREVQIN   RLIRRMN   GTMEGVEEKKKEVPAVPETLK   KKRRNFAELKIKKLRKKFAQK   MKRKARKLIPKKAFAQK   MKRKARKLIPKKAFAQK   MKRKARKLIPKKAFAQK   RQMYRITEIRMANMARKAGN   FYVPARTQIWRFVIRIRGINGS   EPQRFRKVLQLLRPFVKIFNG/T   PFVKLQQGFRNMLRDCRSHIL   AWGIYPNLLSVSPELIYKGRGY   QNQFDLIFFIEKRSHPGSQ   QYPVKDKLLLEVGTGGSFPSW   ULMGOPSSPSGIFGAFQRGLLR   GTYNKYINVKKGDLGGFTW   LACYVLFSYSFSYKHLKHER   LRKYH   LACYVLFSYSFSYKHLKHER   LRKYH   LRYH   LACYVLFSYSFSYKHLKHER   LRKYH   LRYH						1	,
TEIRMARMATKAGNIFYYPAEP   KLAFVIRIRGINGVSPKVRKVI.Q   LFLLYQIFNGAFVKLINKASINM   LRIVEPYIAWRYPNILKSVNELI   YKCGYGKINKK-MAIDNALTA   RSLGKYGIIGREDLIHEIYTVGK   CFKEANNFLWPFI/KJSSPRGGM   KKKTTHEVEGGDAGKREVQIN   RLIRRINN   GTMEOVEEKKEVPAVPETI.K   KKRRNFAELKIKRI.RKKFAQK   MLRKARRKLIYEKAKHYHKEY   RQMYRIVERMAIRMARKAGN   FIVYPARTQIWRFVIRINGGINGS   EPORFRKVJQLIAPPVIRINGGINGS   EPORFRKVJQLIAPPVIRINGGINGS   EPORFRKVJQLIAPPVIRINGGINGS   ONQPEDLHFPTEKRSHPGSQ   ONQPEDLHFPTEKRSHPGSQ   ONQPEDLHFPTEKRSHPGSQ   OYPVKDKKLIEVCYTGGSFPSW   ULMIGDSSPSGIFGAFQRGLIR   GTYNKYINVKGSDLGGFTM   LRKYH							
LRIVEPYIA/WKYPNLKSVNELI							
LFLLYQIFINGAFVKLNKASINM	İ						KLAFVIRIRGINGVSPKVRKVLO
LRIVEPYIA/WKYPNLKSVNELI	ì						
RSLGKYGIIGREDLIHEIYTVGK							
RSLGKYGIIGREDLIHEIYTVGK							YKCGYGKINKK*IALTDNALTA
	1		1		i		CFKEANNFLWPF/K/LSSPRGGM
RLIRRNN							
						ŀ	
	23062	53430	A	23196	3	571	GTMEGVEEKKKEVPAVPETLK
M.R.K.A.R.K.LIYEKA.K.H.Y.H.K.EY RQMYRT\EIRMA\RMARKI.AGN FIYVPARTQIMER\GYIRWARMARKI.AGN FIYVPARTQIMER\GYIRWARMARKI.AGN FIYVPARTQIMER\GYIRWARMARKI.AGN FIYVPARTQIMER\GYIRWARMARKI.AGN FIYVPARTQIMER\GYIRWARGA FIYVPARTQIMER\GYIRWARGA\GYIRWA			-				
FVYPARTQIWREVIRURGINGS   EPQRFRKVLQLLRPFVKIFNG/T					i		
FVYPARTQIWREVIRURGINGS   EPQRFRKVLQLLRPFVKIFNG/T							
PEQRERKVIQLLERPFKIENG/T   PFVKLQQGFRINMLRDCRSHIL   AWGIYPNLKSVNELIVKIRGVG   QNQFDLHFPTEKRSHFGSQ   QNQFDLHFPTEKRSHFGSQ   QNQFDLHFPTEKRSHFGSQ   QNQFDLHFPTEKRSHFGSQ   QYPVKDKKLLEVQTGGSFPSW   ULMKGPSSPSGIFGAFQRGLLR   GTYNKYINNVKGSDLGGFTMV   LACYVLFSYSFSYKHLKHGER   LRKYH   LRKYH   PRVRLRC*MVDEYCGCEWGK*   LNGGRVLWM   LNGGRVLWM   QACIMEPYKLVVLGSGGVGKS   ALTVQFVQGIFVEKYDPTIEDS   YRKQVEVDAQQCMLEILDTAG   TEQFTAMRDLYMKNQGFALV   YSTTAQSTFNDLQDLREQLIRV   KDTDDYPMILVGNKCDLEDER   VVGKEQGQNLARQWNNCAFL   ESSAKSKIVNEIFYDLVRQINR   KTYPYPGKAPQKSPSQCLL   SPHGEPLLGQCRREMWGWSLH   LS3066   53434   A 23200   461   1422   SPHGEPLLGQCRREMWGWSLH   LS3067   53435   A 23201   232   331   TPPL*KWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVYTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQT   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQT   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAVTVTPQTG   TEPL-SWLRGVMAV							
PFVKLQQGFRFNMLRDCRSHIL   AWGIYPNILKSVNELIYK\RGYG   QNQFDLHFPTEKRSHFGSQ   QNQFDLHFPTEKRSHFGSQ   QNQFDLHFPTEKRSHFGSQ   QNQFDLHFPTEKRSHFGSQ   QYPVKDKKLLEVQTGGSFPSW   ULMGDSSPSGIFGAFQRGLLR   GTYNKYINVKKGSDLGGFTMV   LACYVLFSYSFSYKHLKHERI   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRKYH   LRYH							
QNQFDLHFTEKRSHPGSQ			1		1		
23063   33431					1		AWG\YPN\LKSVNELIYK\RGYG
QYPVKDKKLLEVQTGGSFPSW   ULMKGPSSPSGIFGAFQRGLR   GTYNKYNNVKKGSDLGGFTMV   LACYVLFSYSFSYKHLKHŒR   LRKYH   LACYVLFSYSFSYKHLKHŒR   LRKYH   ENGERVLWM   CACKMEPKLVVLGSGGVGKS   LNGGRVLWM   CACKMEPKLVVLGSGGVGKS   ALTVQFVQGIFVEKYDPTIEDS   YRKQVEVDAQQCMLEILDTAG   TEQFTAMRDLYMKNQGFALV   YSTTAQSTFNDLQDLREQLLRV   KDTDDYPMILVGNKCDLEDER   VVGKEQGQNLARQWNNCAFL   ESSAKSKIVNEIFYDLVRQINR   KTYPVPGKAPQKSPSCQLL   SPHGEPLLGQCRREMWGWSLH   CACKMEPKL   CACKM			1				QNQFDLHFPTEKRSHPGSQ
NILMRGPSSPSGIFGAFQRGLLR GTYNKYRNVKKGSDLGGFTMV	23063	53431	Α	23197	2	340	PRVRPRVRDSKMASVGEVSGP
GTYNKYRNVKKGSDLGGFTMV			1				QYPVKDKKLL\EVQTGGSFPSW
LACYVLFSYSFSYKHLKfier			ļ.				\ILMRGPSSPSGIFG\AFQRGLLR
LRKYH			į .		1		GTYNKYI\NVKKGSDLGGFTMV
23064   53432			1				L\ACY\VLFSYSFSYKHLKH\ER\
LNGGRVLWM	1				l		
23065   53433   A   23199   123   693   QACIMREYKLVVLGSGGVGKS   ALTVQEVQGIFVEKYDPTIES   YRKQVEVDAQQGIFVEKYDPTIES   YRKQVEVDAQQGMLEILDTAG   TEQFTAMRDLYMKNGGGFALV   YSITAQSTFNDLQDLREQILRV   KDTDDVPMILVGNKCDLEDER   VYGKEQGQNLARQWNNCAFL   ESSAKSKIYVNEIFYDLVRQINR   KTPVPGGKAPQKSPSCQLL   SPHGEPLLGQCRREMWGWSLH   TESPLGQWGSQILKLQNDLL   23067   53435   A   23201   232   331   TPL*KVLRGVMAVTVVTPQTG	23064	53432	Α	23198	2	48	PRVRLRC*MVDEYCGCEWGK*
ALTVQFVQGIFVEKYDPTIEDS YRKQVEVDAQQCMLEILDTAG TEQFTAMRDLYMKNQGGFALV YSITAQSTFN0LQDLREQILRV KDTDDYPMILVONKCOLEDER VVGKEQGQNLARQWNNCAFL ESSAKSKIYVNEIFYDLVRQINR KTYPYPGKAPQKSPSCQLL 23066 53434 A 23200 461 1422 SPHGEPLLGQCREBMWGWSLH TESPLGQWGSQILKLQNDLL 23067 53435 A 23201 232 331 TPL*KWLRGVMAVTVYTPQTG							
YRKQVEVDAQQCMLEILDTAG   TEQFTAMRDLYMKNQQGFALV   YSTTAQSTFNDLQDLEEQLIRV   KDTDDVPMILVGNKCDLEDER   VVGKEQQQNLARQWNNCAFL   ESSAKSKIVYNEIFYDLVRQINR   KTPIVPGIKAPQKSPSCQLL   Z3066   53434   A   23200   461   1422   SPHGEPLLGQCRREMWGWSLH   TESPLCIQWGSQILKLQNDLL   Z3067   53435   A   23201   232   331   TPI.*KWLRGVMAVTVYTPQTG	23065	53433	Α	23199	123	693	
TEQFTAMRDLYMKNGGGFALV   YSITAQSTFNDLQDLREQILRV   KDTDDVPMILVGNKCDLEDER   VVGKEQGQNLARQWNNCAFL   ESSAKSKIYVNEIFYDLVRQINR   KTPVPGKAPQKSPSCQLL			1	ł	İ		ALTVQFVQGIFVEKYDPTIEDS
VSITAQSTFNIDLQDLREQUIRV   KDTDDVPMILVGNKCDLEDER   VVGKEQGQNLARQWNNCAFL   ESSAKSKIYVNEIFYDLVRQINR   KTPVPGKAPQKSPSQLL   23066   53434   A   23200   461   1422   SPHGEPLLGQCREBMWGWSLH   TIESPLCQWGSQILKLQNDLL   23067   53435   A   23201   232   331   TPL*KWLRGVMAVTVYTPQTG	1		1				
RDTDDVPMILVGNKCDLEDER   VVGKEQQNLARQWNNCAFL   ESSAKSKIYVNEIFYDLVRQINR   KTPIVPGIKAPQKSPSCQLL   23066   53434   A   23200   461   1422   SPHGEPLLGQCRREMWGWSLH   TESPLCIQWGSQILKLQNDLL   23067   53435   A   23201   232   331   TPL*KWLRGVMAVTVYTPQTG	1		1		ł		TEQFTAMRDLYMKNGQGFALV
VVGKEQGNLARQWNNCAFL   ESSAKSKIYVNEIFYDLVRQINR   KTPVPG0KAPQKSPSCQLL   23066   53434   A   23200   461   1422   SPHGEPLLGQCRREMWGWSLH   TESPLGQWGSQILKLQNDLL   23067   53435   A   23201   232   331   TPL*KWLRGVMAVTVYTPGTG			1		1		
ESSAKSKIYVNEIFYDLVRQINR   KTPVPGKAPQKSPSCQLL	1					1	
KTP\VPG\KAPQKSPSCQLL	1			l	1		VVGKEQGQNLARQWNNCAFL
23066         53434         A         23200         461         I422         SPHGEPLIGOCREMWGWSLH           23067         53435         A         23201         232         331         TPL*KWLRGVMAVTVYTPQTG						l	
TESPLG\QWGSQILKLQNDLL 23067 53435 A 23201 232 331 TPL*KWLRGVMAVTVVTPQTG			L				
23067 53435 A 23201 232 331 TPL*KWLRGVMAVTVVTPQTG	23066	53434	A	23200	461	1422	
			L				
I I I I I I I I I I I I I I I I I I I	23067	53435	Α	23201	232	331	TPL*KWLRGVMAVTVVTPQTG
IIAII DI MILWOSTOTI KETA			L				IIAIPDPHHEWGSTGYPRLPA

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23068	53436	A	23202	5	1223	I GLDSGPESGLRAQPGLLAGKRK
23008	133430	^	23202	ľ	1223	MSVLRRMMRVSNRSLLAFIFFF
	1				1	SLSSSCLYFIYVAPGIANTYLFM
	1	1				VOARGIMLRENVKTIGHMIRLY
		l				TNKNSTLNGTDYPEGNNSSDYL
		1				VOTTTYLPENFTYSPYLPCPEKL
		1				PYMRGFLNVNVSEVSFDEIHOL
		1				FSKDLDIEPGGHWRPKDCKPR
		1				WKVAVLIPFRNRHEHLPIFFLHL
		1				IPMLQKQRLEFAFYVVEQTGTQ
					1	PFNRAMLFNVGFKEAMKDSV
						WDCVIFHDVDHLPENDRNYYG
1		ŀ				CGEMPRHFAAKLDKYMYILPY
						KEFFGGVSGLTVEQFRKINGFP NAFWG\WGGEDDDLW\NRV\H
		1				YAGY\NV\TRPEGDLGKYKSIPH
					ŀ	HHRGEVQFLGRYKLLRYSKER
		1				QYIDGLNNLIYRPKILVDRLYT
		1				NISVNLMPELAPIEDY
23069	53437	A	23203	78	495	CTFEKAPPFOLGRGRMVPAKK
23009	33437	^	23203	/°	493	GGEKKTGR\AINEVVTREYTINI
		1				HKRIHGVGFKKRAPWALKEIR
		1				KFAMKEMGTPDVRIDTRLNKA
		1	ł			VWAKGIRNVLYRIHVRLSRKR
		1				NEDEDSPNKLYTLVTYVTVTTF
		ı			ì	KNLOTVNVDEN
23070	53438	A	23204	138	360	KNOKROLSGGHCADOTSLVLA
23070	33436	<u> </u> ^	23204	138	500	RLAPLF\PTTMSDKPDMAEIEKF
	1	ı				DKSKLKKTETQEKNPLPSKETT
		1				EQEKQAGKS
23071	53439	A	23205	145	728	KRQPTSAMKDPSRSSTSPSIINE
25071	33437	ľ.	20200	145	/20	DVIINGHSHEDDNPFA\EYMWD
1	ŀ					GKMEGRNFNR\OIGRRELWGR
1		1				KD\FLNRCFPKE\MLGEEGKEG
	1					ALNWVLFPSFEILPHOTMGPNP
	İ	l				RTSFNDPCLSSDGSFSWEDLVV
	į.		i	ļ		KS\NLNPNAKEFVPGGEVRKYL
	l					SRRGPLLVDVAQFPHCEGSIRR
		1		i		LNCKSSLGHCVTLYAIAQSFC
23072	53440	A	23206	530	707	Quic
23073	53441	A	23207	1	515	MDFSKVPPHAAFRAQPPPALT
[		ľ	l	ľ		WGLAKGSGRGAGEGGERPDRR
						R/PAARTGTPAWRPAGRRRLPP
	1	l	1	I		RSPPPAPVPVRPALPHPVRPGKE
	1	l	1	l		TGKTRRRRHYSPGRPLRGAAA
	1	l		l		AAAAAAVAOTASPSWAEGGRS
1	1	1		I		GAGGSAESPAFSSPALPCPEGLL
				1		CPSAVPVPSKQSVSEVLAIF
23074	53442	A	23208	3	713	C.C TIGINGO TOD TOAIL
23075	53443	Ā	23209	1	704	
23073	1222	٠.	20207	ı	/ / .	

	01/0/306/					PC1/US01/08631
SEQ ID	SEQ ID NO:					Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	l	09/540,217	sequence	or peptide sequence	deletion, (-possible nucleotide insertion)
				o-quemee		
23076	53444	A	23210	2	1631	GLPRRPTRPPTRSGVSPPWPGPF
						SGVLFPLLLHCTGSLSQAVLTQ
		l			1	PSSLSASPG\ASVKLTCTLRSDIN
						VGSPFIYWYQQKPGSPPQFLLR
		l				YKSDSDNQQGSGVPSRFSGSKD
		Į			i	ASANAGILLISGLQSE\VRDDYY
		1				CQAWDSSTAHSDTGRCGIVGRI
		1				PVRQCWPLLLLGLAMVAHGLL
						RPMVAPQSGDPDPGASVGSSRS
		İ		1		SLRSLWGRLLLQPSPQRADPRC
						WPRGFWSEPQSLCYVFGTGTK
						VTVLGOPKANPTVTLFPPSSEEL
						OANKATLVCLISDFYPGAVTVA
						WKADGSPVKAGVETTKPSKOS
1						NNKYAASSYLSLTPEQWKSHR
						SYSCOLOWGLGLGASQGGFLY
						EPVSQCVVFGGGTKLTVLGQP
						KAAPSVTLFPPSSEELQANKAT
						LVCLISDFYPGAVTVAWKADSS
1						PVKAGVETTTPSKQSNNKYAA
1						SSYLSLTPEQWKSHRSYSCQKP
						NHASHGTSRMPSVSAPVSGLKS
		1				CWSGTGDRSALGQMPGPCHGI
1						PDPMQQASDSGQPKAAPLGSL
22000	53445	ļ	22211	1	959	CSKPSSEELSGQQGPHWCVS
23077	33443	Α	23211	[1	959	VFLLLYLVAMPDPAKSAPAPK
1		ŀ				KGSKKAVTKAQKKDGKKRKR
						SRKESYSVYVYKVLKQVHPDT
1						GISS\KAMGIMNSFVNDIFERI\A
1		ļ			1	GEASRLAHYNKRSTITSREIQTA
]						VRLLLPGELAKHAVFIENLLNY
1						RKYLVKSSEKDRNYPCPLPGLK
1						SKVDICTPYCLLLLPQPRHYPG
			ĺ			KDTKEETEDFPNSPPPINREEDK
						RYTTVMGPCLRQAALEGELLA
						CPVMEDQQANWVHEPITFNTF
						KEIRKSIRENGPASPFTRGLIEVI
1				l		ADSSDLVASTQKLTQPKETPSA
	1					PYNDISDLTNQDSWFIGFPPIWN
						AGETDLSNNKTT
23078	53446	A	23212	3	463	
23079	53447	A	23213	74	464	
23080	53448	Α	23214	41	490	
23081	53449	Α	23215	2	369	PDPAANAPAAAKPAAAKALPV
			1	l		FLLLLSGGAGPAHPMRAPGC*G
1	l					ELGYTEQLFQFLQATAQEEGIF
1	l	İ				EYTDPKLAISAVWTFRDLGVLQ
1	l	l				QTPSPAGPRLHLSPTFASLDNQE
						KLEQFIRQFICS
23082	53450	A	23216	124	301	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nuclcotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide scquence	delction, \=possible nucleotide insertion)
				sequence		
23083	53451	A	23217	3	2426	PGVLSTATFRESGLSPCSKLKHP
						GPRGRLOYSPPTAGOGRLENG
						AGHPOSTGPVATLSVAEYGTO
		l				GGASGPARIPIGSRGCLLAGRA
						EASLWSSGFGMKLEAVTPFLG
			l			KYRPFVGRCCQTCTPKSWESLF
						HRSITDLGFCNVILVKEENTRLF
			ŀ			SWALLRVLKLLFLNVQLHKGQ
						MKMVQKAAQQVGQVGTSLWL
						SPEOGLPLVLLSTHKTLLDGILL
			ŀ			PFMLLSOGLGVLRVAWDSRAC
			l			SPALRALLRKLGGLFLPPEASLS
						LDSSEGLLARAVVOAVIEOLLV
		ļ	ŀ			SGOPLLIFLEEPPGALGPRLSAL
		1				GOAWVGFVVQAVOVGIVPDAL
		l				LVPVAVTYDLVPDAPCDIDHAS
		l				APLGLWTGALAVLRSLWSRWG
		i i				CSHRICSRVHLAOPFSLOEYIVS
		l l				ARSCWGGRQTLEQLLQPIVLGQ
		1		İ		CTAVPDTEKEOEWTPITGPLLA
		l l				LKEEDOLLVRRLSCHVLSASVG
		l				SSAVMSTAIMATLLLFKHOKG
		l				VFLSQLLGEFSWLTEEILVRGFD
		l				VGFSGOLRSLLOHSLSLLRAHV
		l				ALLRIROGDLLVVPQPGPGLTH
		l				LAOLSAELLPVFLSEAVGACAV
1		l				RGLLAGRVPPOGPWELOGILLL
		l				SQNELYRQILLLMHLLPQDLLL
1		l				LKPCOSSYCYCOEVLDRLIOCG
		l				LLVAEETPGSRPACDTGRORLS
		l				RKLLWKPSGDFTDSDSDDFGE
		l				ADGRYFRLSQQSHCPDFFLFLC
23084	53452	A	23218	169	955	ASSQLPCSMAQCCATWAPLKM
		l				RWLCSTAWSTSSLSLPPSWSSL
		l				QASQPSCCHATSLAPPCAGQCL
		l	i			ARAWPVLSFLCTCALGLLASIA
		l				MTFATOGKALLAACTFGSSELL
		l				ALAPYCPFDPTRIYSSSLCLWGI
		l				ALRCSAWAENVFAVRCAOLTH
		1	l	l		QLLELRP\WWGKKQPTHMMRE
			l	l		NPELVEG\RDLLSCT\SSE\PLTL
		l		1		CRDDVLPRP\DGQLRTLQATLL
				l		CDPGPGFLGAGLRGLKWTLGG
		l	l	l		P\SGAFQPSSPTTPAHLHLKIETL
		l				F
23085	53453	A	23219	h	344	
25005	100.00	٠.	22217	L	2	

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	l	02/340,217	sequence	or peptide sequence	deletion, (-possible nucleotide insertion)
		1				
23086	53454	Α	23220	214	1355	FCSSWLRSLADSSLSWKMFLVC
	ł	1				LTGGIASGKSSVIQVFQQLGCA
	ŀ	1				VIDVDVTARHVVQPGYPAHRRI
	ŀ	l				VEVFGTEVLLENGDINRKVLGD
	ľ	l				LIFNQPDRRES\LNAITHPEIRKE
	Į.	l		ì		MMKETFKYFLRGYRYVILDIPL
		l		i .		LFETKKLLKYMKHTVVVYCDR
	1	1				DTQLARLMRRNSLNRKDAEAR
	1	l				INAQLSLTDKARMARHVLDNS
	1	l		ì		GEWSVTKRQVILLHTELERSLE
		l				YLPLRFGVLTGLAAIASLLYLLT
		l				HYLLPYAYTASAIPPSGAASRG
		l		1		PGARCSAIRKRAPARPTSAVSPP
	ŀ	l		l .		GAPRSPARGLRAIDRAAGTRNL
		l				NATLPPPGPPAPPSPPSPPSSITL
	l .	l				HVFHQLQPDITRGREGTDFQWE
				l		KFPGRVLVADSRRALRCLAPAP
				·		TWLV
23087	53455	Λ	23221	2	1231	TPLFLRKGTADGSARTTVIHLG
		ł	:	l		FTKGKQRCVETQAAAVAGDVG
	ì	l	1	l		RTAAGGRVGKESGVISSDDRPG
	l	1	İ			LPVGQSQPEGGEQEATDTVSVH
		l	1	l		ATTSQAWPGGADSKELSSKAE
		1		ŀ		ERETGKDQQTELPLCLPLRDFP
					l .	DSAPVMSSDSGHATESLHLASS
				l		DSKPPMPTPNEGPGSTVGHAGP
				l	l	PRTVKADLCTEPQLALRGLQVS
					1	LPRLHLVGVIYAEHQASGLEGC
				l		GHAGSCSPDGSPRAPGATLCLS
				l	1	KPLSFRTKRCLPCWLPASAPVGI
				i		PGAPVHIPGAPVCIPGAPACISN
		l		l .		APVHIPGAPVGIPGAPVCIPGAA
		1	İ	l		VCVPNTPMCIPSAPMCIPNTPVC
		1	1	l	1	IPNTPVCIPNTPVCIPGAAVCILG
		l	I	l		A/PRAHPWCSSCVHPGAPECYL
		l		ŀ		GCGLTQADRLEPAVLDAGLQE
		<u></u>				PLFALCGQALPS
23088	53456	Α	23222	2	408	
23089	53457	A	23223	62	251	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence	İ	
		<u> </u>				
23090	53458	A	23224	2111	3003	TQKAICMCLGLCKKIHHLLSLQ
1	l					CLISQIKMIMGAGWQ/SSGFTDE
1	l	1				KETSKIKADPEKLKKLQESLLE
1						EDSEEEGDLCRICQIAGGSPSNP
1	1	1				LLEPCGCVGSLQFVHQECLKK
		l			Į.	WLKVKITSGADLGAVKTCEMC
l					i e	KQGLLVDLGDFNMIEFYQKHQ
		1				QSQIKTEATVVEHKDIYNVTET
						QLVPEARSVLLVGAGQMLLTG
i						ANRALNSDYSSMRPKVKAQNE
					İ	LMNSGLYLVLLLHLYEQRFAEL
		l				MRLNHNOVERERVRVTRVLGG
1		1				LSPNVSCGVVSGVGSFQWVRG
		1				LTDLKNEASDLHAQIPAE
23091	53459	A	23225	ī	594	MIAAFNPGKVDTVTTNGPFIDL
		1				NYMVYMFHYDPTHGKFHDVP
1		l				MFVMGMNHKHENKLTVISNAS
		1	i			CTTNCLDPLAKVIHDNFGIREG
1		1				LMTTVHSIIATQKTVDGPSRKL
					Į.	WHDSHGAFONTIPASTGAAK\I
l		l				KKVVRQA/SQGPLKGILGYTEH
		1				OVVPLEFKSNTRPSTFEAGAGI
				i		AVNDHFVKLISWYGNEFAYSN
						KVIKPM
23092	53460	Α	23226	3	232	
23093	53461	Α	23227	1078	2221	QQTLQPAALPCPFPPPTHPPEVL
					1	PTAMNTTSSAAPPSLGVEFISLL
	l					AIILLSVALAVGLPGNSFVVWSI
l					ŀ	LKRMQKRSVTALMVLNLALAD
						LAVLLTAPFFLHFLAQGTWSFG
						LAGCRLCHYVCGVSMYASVLL
						ITAMSLDRSLAVARPFVSQKLR
						TKAMARRVLAGIWVLSFLLAT
i					ŀ	PVLAYRTVVPWKTNMSLCFPR
						YPSEGHRAFHLIFEAVTGFLLPF
						LAVVASYSDIGRRLQARRFRRS
		1				RRTGRLVVLIILTFAAFWLPYH
1		1		1		VVNLAEAGRALAGQAAGLGLV
1		1	l	I	ŀ	GKRLSLARNVLIALAFLSSSVNP
1		1	1	ŀ		
l		1		1		VL\YACAGGG\LLRSAGVGFF\A
				l		KLLEG\TGSEASSTP\RGGSLGQ
l			l			TV\KSGPAALEPGPSESLTASSP
		١.	00000	6105	(212	LKL\NELN
23094	53462	A	23228	6125	6319	ASVVPVTSALAMCSPKSGIGDG
l	1	ı	l	1	l	FAVLRVEDVAQK***KAALGV
22225	52462	١	2222		1242	ASSPA WQQTREREHVQGNSPL
23095	53463	Α	23229	I	1347	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23096	53464	A	23230	1	1124	MRECISIHVGQAGVQIGNACWE LYCLEHGIQPDGQMPSDKTIGG
		1	İ			GDDSFNTFFSETGAGKHVPRAV
		1				FVDLEPTVVDEVRTGTYRQLFH
		1				PEOLITGKEDAASNYARGHYTI
1		1				GKEIVDLVLDRIRKL\FGGGTGS
1		l	l			GFASLLMERLSVDYSKKSKLEF
		1	ŀ		Ì	AIYP\APQVSTAVVEPYNSILTT
						HTTLEHSDCAFMVDNEAIYDIC
		1				RRNLDIERPTYTNLNRLIGQIVS
1		1				SITASLRFDGALNVDLTEFQTN
						LVPYPRIHFPLATYAPVISAEKA
ł						YHEQLSVAEITNACFEPANQMV
		1				KCDPRHGKYMACCMLY\RGDV
				i .		\VPKDVNAAIATIKTKRTIQFVD
1						WCPTGFKV*LGDMEAFQQAAD
1						AQNNTGLEGPHPLGRLSLFHGL
23097	53465	Α	2323 I	I	2918	MDVYCAHESLRHTKSPEATGQ
		i				TSGQRPEALQAFWPAFGSSPVT
1		i .				GSSPVTEVTTKGRWSSSRQVL
İ		1				WDRIPLAKSRPAISNRQSQRTNI
		1				PWWESCPPILNLEREQPPRFAQP
		l				GTYASRWKALEEMEKQQREQ
		1				VDRNIREAKEKLEAEMEAARH
		1				EHQLMLMRQDPGLALVSCMQ
	İ				İ	QGQSMSRSPKKVFKHLYELNL
					Ì	TKLQSLGSRSRLRWKDSPFWGE
		į .	1		I	EAGGKRPPLVSLALHRSPSLLL
22000	60466	<u> </u>	22222	,	158	LFVPKHPSKGRPDCTVQRS
23098	53466	A	23232	1	414	LGGLKGTWGETKPAMAAAEEE
23099	33407	^	23233	'	414	DGGPEGPNRERGGAGATKTLL
		1				YTGNHLFSPOWLETRPEROECP
					1	VCKAGISREKVVGEKSLLTFSL
						HFLRLKTPOICRLPV*LFLPP*GF
						OPFGDTGGFHFSFGVGAFPFGF
		1				FNSPSY
23100	53468	A	23234	1	663	
23101	53469	A	23235	2	667	SWAGGGFWFVGACVCVFGGTE
		1		[ _		GYVGRNKTGHGSSRRRRSGAP
		1				NVPNRERGGAGRPKTPFECNIC
		1	1			LETARELVLSVLANLYFWTISH
					1	QV\LE\TRPE\RQECP\VCKAGISR
		1			I	EKVVP\LYGRGSQEAPRIPRLKT
		1			1	PPPPPRGQR\PAPE\SRGGF\QPFG
	1	1			1	DTGGFHF\SFGVGAFPFGFFTTV
					1	F\NAHEPFRRGTGVDLG\QGHP
					1	A\SSW\QD\SLLLFLAIFFFF\WLL
23102	53470	A	23236	363	476	
23103	53471	В	23237	224	364	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \"possible nucleotide insertion)
				sequence		
23104	53472	Α	23238	1	1324	MRQRRAELLSKDKRVRTDGNN
						KCSPSAIGKRKERGTNDWIEGD
		l				NNQESQGLNCSDTLNRDLGPN
i		l		i		TRGFLYAGLSGLDPDPSLPTPD
		1				MSSEVLEDNLDTLSLYSGKDSD
						SVKLLEEYADSESQASLQGRSR
						CWQHPHPAPLSYPPAASGSQLS
						LQGKFPSAKPPLLGGYTCTAQR
				l		TQWALTKRSLRATRESVSRQQ
						GVYGLVGRKRTRHHRGVLGLL
		1				GIPSARPERPGTGSEAGPTRLPG
						AAVARRRPARPVRGRLGDPAP
1						TAAQTSGPASLPPPPPPRSYGDP
						HONGYPDPDPAPSASPDLPPAA
					l	GDTRGSGGRRKKRRTGTARLN
						TPGRKA\PKQKAAAGRKGRAH
						OPAPRTPEPORRRPIRPPNOTOR
	i					LRHPHRRAGRGPPPGGAGGTA
						NPSTGANGGAATARRTAGART
						RGARGASERAGRHGSRPASWL
	İ					TATPPPGT
23105	53473	A	23239	1	433	PTRPSSPEETDGPWDPEGHPGG
25105	33473	ľ	23237	ľ	455	PLTDPPNAPVLALPPAIPGLSLS
		1				MADGSSDAAREPRPAPAPIRRR
						SSNYRAYATELODLCROLHAR
					i	VDKVDEERYDIEAK\VTKNITEI
						ADLTQKIFDLRGKFKRPTLRRV
						RISADAMMQALL
23106	53474	Α	23240	2	409	RISADAMINQAEE
23100	53475	Ā	23240	3	490	SSPEETVGPWDPEGHPGGPLTD
23107	33473	<u> </u> ^	23241	ľ	450	PPNAPVLALPPAIPGLSLIMADG
		1				SSDAVRAAG*GVAGTPRARAR
					1	EPRPAPAPIRRRSSNYRAYATEP
		1				HAKKKSKISASRKLOLKTLLLO
	1	Į.				IAKOELEREAEERRGEKGRALS
		İ	į			TRCQPLELAGLGFAELQDLCRQ
i		1			1	
	******	<u>.                                    </u>	02242	1.40	046	LHARVDK
23108	53476	Α	23242	149	846	VSAWRNGSSDAAREPRPAPAPI
1		l		I	1	RRRSSNYRAYATEPHAKKKSKI
		1		I		S\ASRKLQLKTLLLQIAKQE\LE
		1			1	REAEERRGREGARSLAPRC\QP\
		l	1	1	1	LELAG\LGFADLPGLCCRQFHA
1		l				RV\DKVDEERYDIEAKVSKNITE
	1				1	IAN\LT\QKIFDLRSKFKRPTL\RR
		l	1	1	I	VRISADAMMQALLGARAKESL
	1				1	DLRAHLKQVKKEDPEKENREV
	1	1		1	1	GDWRKNIDALSGMEGRMKKW
1						
1						RELSLPAYCPCPEEGH
23109	53477	A	23243 23244	1 80	487	RELSLPA YCPCPEEGH

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, ^=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23111	53479	A	23245	427	1417	KTMWKVEFGVRFSITRLLEHQ PKQTMYIPLPK/PPLWPEEPKTE APKQTPRAPPKPKTSPRPRIPQT QPEETDQSTQEPFTTKIPRTTEL
}	İ					AKTTQGLEMLSCLIHNPKMHK
						DSTNEQIAASLQACYYEQLLM
						YIRVTDVDSVGPGNGKEAVTA
ŀ						PQAPTLFWLNHKAQLAKDWA
						GKDVNITPTFATSKKENKPPNN
						LQICNVSKIHQRAKVAGKPEIK
						DKHIQKEMPLDAIEASKINLVKI
						FNEFLKAWKCRLSFEYRIPENQ
						GHKGSFYLLVGSSRTSWLLTEK
		1				IGSSKRVLRMPPMDILKTTQILL
		_				SFLWSINLTPRGKNNHLVAPAH
23112	53480	Α	23246	3	1011	
23113	53481	A	23247	2	334	PIQSFFLVLKKPPLRKGVYLSSK SLSRWFFKVQAN*RSQPQKAFS RLSCHC
23114	53482	١.	23248	1	642	RESCRE
23114	53482	A A	23248	854	1150	KEDTNKWKNIPCSWLGRINIVK
23113	33463	^	23249	0.54	1130	MAIMPKVNL*IOCHPHOATNDF
ì		l				SSQNWKKTTLKFIWEPKKEPAL
		l				PSQS*AKRAKLGGITLPDFQTIT
1		l				YKATVTKTA
23116	53484	A	23250	1754	2393	PSNRREFGESWPETQAPGALRP
23110	33404	ľ`	23230	1,754	12373	LLPAPGTPRARERPAGWPARPM
	ľ	1				EVYIPSFRYEESDLERGYTVFKI
ŀ		1				EVLMNGRKHFVEKRYSEFHAL
İ		1				HKKLKKCIKTPEIPSKHVRNWV
i		l				PKVLEQRRQGLETYLQ\AVILE
1		1				NEELPKLFLDFLNVRHLPSLPK
		l				AESCGSFDETESEESSKLSHQPV
		1				LLFLRDPYVLPAASDFPNVVIE
1		1				GVLHGIFYPHLQPR
23117	53485	A	23251	1	265	AGNRLRFQLELEFVQCLANPNY
		1				LN/WYPQCLHMLELLQYEHFR
		1				KELVNAQCAKFIDEQQILHWQ
1		l				HYSRKRMRLQQAL\AKQQQQN
		1				NTSGK
23118	53486	Α	23252	1	987	
23119	53487	Α	23253	1	491	RGFRNFARVSGLLLCQAGGVL
l l		1			l	VSSFVMAAAVAMETDDAGNR
		1				LRFQLELEFVQCYANHNTLIFLP
1		1			1	QR\GYF\KDKAFVNYLKYLLYW
1					1	\KDPEYAKYLKYPQCLHMLGA
1		1	l	1	1	APNMEHFRKEAGWNAQC\AKF
						IGWNRQVSTLGKHYS\RKRM\R
L			L			LQQALAEQQQQNNTSGK

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, /=possible nucleotide insertion)
23120	53488	A	23254	73	355	HEGOVHVLLASAHVDVAIVLG
23120	33400	1	23234	1,3	333	AGSMVEGEARPEOAAOHHOGI
		1				APLHLDAPGPSWTOSRSPRNRA
		1				NONHTCSEAT*SHPEOLPOTLW
		ı				ECHAPRSP
23121	53489	В	23255	1	441	
23122	53490	Ā	23256	528	698	HGQRFLHGPV/H/HHSHYDN/PH
						HSLHHHRCLHHCHLHHHRHHL
						HHHPLHCYHHHPGQNG
23123	53491	Α	23257	1	403	
23124	53492	Α	23258	509	4547	LYLTTSRREEHLKSLVVETQDL
						ASPVLRSVSICTKVEEAFRQAH
1						VIVVLDDSTNKEVFTLEDCLRS
						RVPLCRLYGYLIEKNAHESVRV
		1				IVGGRTFVNLKTVLLMRYAPRI
1		1	İ			AHNIIAVALGVEGEAKAILARK
						LKTAPSYIKDVIIWGNISGNNYV
1				٠.	•	DLRKTRVYRYESAIWGPLHYSR
					ľ	PVLNLIFDSEWVKREFVAILKN
						LTTTGRQFGGILAAHSIATTLKY
ł						WYHGSPPGEIVSLGILSEGQFGI
						PKGIVFSMPV
23125	53493	A	23259	56	290	
23126	53494	A	23260	468	993	VLLACFWPGSDFWPRSRRKTY
					1	GTAPQSCYSFYVALDCGGVSRL
l	İ					VIFVSWRNPQVAPTSAHQNRPS
f	ł				i	RNPVSRPPNTQRVARRKHYAL
ľ	1		l	l		ADGYTERRWTNAPCRAESSFPP
ŀ			l			NCPSAAPIPDSYDK*PSRATLFT
						SHLTVEAFPVWSYLYPGETPRL
ľ						HPRAHTRTVPAAIRYPDRQTPN
				i		GLRGANTMHSQMDTLNGDGR
ŀ						THLAGQRARSPRTARPQRQYRI
1						HTTSDLPGLVRKYASGCCLRIW
İ		1				YCCAWIERALSLQVGGGVFGD
23127	53495	-	23261	1	3114	PLSEGQAECDHGST
23127	53495	A	23262	3	401	TSSLSLSGKSGRYIVFLRRSVGI
23126	23490	<u>۱</u> ^	2.3202	2	401	OSPSAVATVRLLLAGSDRRFAA
1		l	1			GSAGCAVLSRAERS*EPGCCYR
1	l	İ				IRRSAFRLRAGSIGNRDKPVAW
	l	l	I			PDTGSGDNTEVYFRAHQAGILP
l		l				DTVLAGPAQRVAVSGGTPVDW
23129	53497	A	23263	2874	3062	DI VEAGI AQRVAVSOGIT VDW
23129	2347/	<u></u>	23203	2014	3002	L_,,

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23130	53498	A	23264	2766	3323	MPDLVEGREKSDKGAQPQPRT OSRSARTNGSGVRILHSLWRIIG IPLLLGYSLVCSRVLLACFWPG SDFWPRSRRKTYGTAPQSCYSF YVALDCGGVSRLVIFVSWRNP QVAPTSAHQNRPSRNPVSRPPN TORVARRKHYALADGYTERR WTDAPCRAESSFPPNCPL*RQY RIHIVDASVCGP
23131	53499	Α	23265	968	1440	
23132	53500	A	23266	3	640	LSVVRSRIHMGVGVGCPVR YRNQDDHELQTHGNKIL.CGIV CDKGAQPQPRTQSRSARTNGSG VRILHSLWRIIGIPLLLGYSLVCS RVLLACFWGSDFWPRSRRKT YGTVPOSCYSFYVALDCGGVS RLVIFVSWRNPQVAPTSAHQNR PRRNPVSRPPNTQRVARRKHYA PADGYTERRWTIAPCRAESSFP PNCPSAAPIPDSYDK
23133	53501	A	23267		3140	MVYKMRYRSOHPYSIKEKOMK SEVLSVKEKIGYGMGDAASHII FDNVMLYMMFFYTDIFGIPAGF VGTMFLVARALDAISDPCMGL LADRTRSRWGKFRPWVLFGAL PGIVCVLAYSTPDLSNNGKMI YAAITYTLLTLLYTVVNIPYCA LGGVITNDPTQRISLQSWRFVL ATAGGMLSTVLMMPLVYLIGG DNXPLGFGGDLSPMFSTPEEIA RCPPYENDVHVGASSLAAGH KTLIPELVRSAEQHMGTR
23134	53502	A	23268	750	1340	ILHAPAPPESASASHEQFEWSD KGAQPOPRTQSRSARTNGSGVR ILHASLWRIGIPLLLGYSLVCSRV LLACFWPGSDFWPRSRRKTYG TAPQSCYSFVVALDCGGVSRLV IEVSWRNPQVAPTSAHONRPSR NPVSRPPNTQRVARKHYALA DGYTERRWTNAPCRAESSFPPN CPFAAPIPDSYDK-PFRIVS
23135	53503	A	23269	93	699	RASVQOKKLSTIDKGAOPOPR TOSSRARTNOSGVRILHSI, WRII GIPLLLGYSLVCSRVLLACFWP GSDFWBRSRKRTYGTAPOSCVS FVYALDCGGVSRKRSYLYPOET FRVAPTSAHONRPSKROTTRPP NTORVARRKHYALADGYTER SATNAPCRAESSFPPELPVPQVP NTGFIRQVTFRDCFVNTPQAAA SGFW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23136	53504	Α	23270	1000	1530	QRYSDNPPNDKGAQPQPRTQSR
						SARTNGSGVRILHSLWRIIGIPL
	ł					LLGYSLVCSRVLLACFWPGSDF
						WPRSRRKTYGTAPQSCYSFYV
		l				ALDCGGVSRLVIFVSWRNPQV
1		l				APTSAHQNRPSRNPVSRPPNTQ
				i		RVARRKHYALADGYTERRWT
		ı				NAPCRA/VEPFPPNCPWEAQYRI
23137	53505	Α	23271	242	711	LKQKRDKEDRSSRVSAVLLIK/
						HRRPLPLPIGLKACHCSCTAKC
l	1					LGGCESPRKGGQLLIFEGHVKI
ŀ		l				DLEIFTKSLWLTRQLIVKKETEK
1						AVKHASNICIMRTGLTFQSHKIE
		l				DGLFGLLVIATEAEWSYVCOST
l	ŀ	l				NPVDVYPLTAVPACCIISASDVP
23138	53506	A	23272	2	459	CVNSOIGRGHKITTYPFSPRKSG
		l				RKGMAOSOGWVKRYIKAFCK
	l	l				GFFVAVPVAVTFLDRVACVAR
l		l				VEGASMOPSLNPGGSQSSDVVL
ŀ		1				LNHWKVRNFEVHRGDIVSLVS
l		1	1			AKNPEOKIIKRVIALEGDIVRFP
	İ	l				*DFCMPMPHISCGPQSAGRNW
23139	53507	A	23273	1	474	
23140	53508	Α	23274	358	948	PLTPFSPRKSGRKGMAQSQGW
1		1				VKRYIKAFCKGFFVAVPVAVTL
	1		1			LDRVACVARVEGASMQPSLNP
1	1	l				GGEPV/ASDVVLLNHWKVTEF/
	Ì	l				SEVHRGDIVFIG\APKNPEQKIIK
i		l				RVICSLKGDIFQNHRTQKP\DVK
l	1	l			1	VPRGSHLGLEGDHSWDTVFDS
1	ı	1				NSFGPVSLGLLHAHATHILWPP
l	1	l				ERWQKLESVLPPERLPGTERRG
		l				MTA
23141	53509	Α	23275	90	664	NKYRLQPSPPRLALWDRASDPP
ŀ						APPEDPETAPAMEEDQELERKI
1						SGLKT\SMVEG\ERKTT\LELV\Q
		l		l		AAG\TDRP\CVTFVLHEEDHTLG
	1			l .		NFLRY\MIMKNP/EKVEFCGYT
	l			i		YG/HPSFQRAKLNLPHSRFRGYP
	ŀ					FPAVEPISREALNELM\NVC\QP
	ŀ					CAFTSFRASIKDYKESKSKQEM
	i	ŀ		1		ESTFLVLLWOFYKGGTVL
23142	53510	С	23276	108	320	` ` ` ` ` `
23143	53511	В	23277	446	625	
23144	53512	A	23278	1	3042	
23145	53513	A	23279	1	3156	
23145	53513	A_	23279	[1	3156	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon,/=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		Ì		sequence		
23146	53514	Α	23280	518	2076	QQRR*MCQNLMTH\QNQRSGR
		1				SQNDF*RRRKNISERSAVYR*R
1		1			İ	QLYIRRQPYAERPS\HVEDKGH
		l				KYLVFEANTGTENGYQGEESLF
		ı				NKAYYGGGTNFFRKESQKLQQ
		l				SAKKRDAELANGALGIIELNND
1		!				YTLKKVMKPLITSNTVTDEIER
1		1			ł	ANVFKMNGKWYLFTDSRGSK
1		ı				MTIDGINSNDIYMLGYVSNSLT
		1				GPYKPLNKTGLVLQMGLDPND
1		l			l	VTFTYSHFAVPQAKGNNVVITS
		İ				YMTNRGFFEDKKATFAPSFLM
		l				NIKGNKTSVVKNSILEQGQLTW
		1				LOVAKRAGLGGGGSGRTVLRE
		l		1		RVRIEIASTHIALAVECVSVRV
1		l				WKVPRLPSRQKYAKHASQLVS
ł				-		NQKSEEEEYMKTFNISQQDLEL
		1				VEVATEKITMLYEDNKHHVGA
		l				AIRTKTGEIISAVHIEAYIGRVTV
1		1				CAEAIAIGSAVSNGOKDFDTIV
į.		1	1			AVRHPYSDEVDRSIRVIPTYGT
İ		1				DEWEOWWNAFNEENLFCSEE
		l	1		1	MPSSDDEATADSQHSTPPKKKR
		1	}			KYIYVLVIAITMVMRFWQYING
23147	53515	A	23281	299	457	KTITYEVIJAINIVIJAA WQTING
23148	53516	A	23282	584	1185	QQRR*MCQNLMTH\QNQRSGR
231.0	55510					SONDF*RRRKNISERSAVYR*R
		1	1			QLYIRRQPYAERPS\HVEDKGH
		į .				KYLVFEANTGTENGYQGEESLF
1		l		1		NKAYYGGGTNFFRKESOKLOO
		l				SAKKRDAELANGALGIIELNND
1		1				YTSEKSNEAADHFKHVPOAKG
		1		l		NNVVITSYMTNRGFFEDKKATF
1		1				APSFLMNIKGNKTSVVKNSILE
		l				
23149	53517	-	23283		3618	QGQLTVN
23149	53517	A	23283	191	819	VALVECCED DICCTODUL FOCI *
23 130	33316	Ι^	23204	191	913	VNVTSFGSRPICSTSPVLFSGL*
1		l	1	l		GPVKEFDT*PSM*ISFELIPSIVIF
		1	1	I		EPRESVNKYHLPFILKTFARSISS
1		1	1	l		VTVFEVISGFITFFNV*SLFNSMI
				l		PRAPFANSASRFLALC*SFWLSL
1					ı	RKKFVPPP*YALLNKDSSPW/SS
			1		1	
						VFCSRVGFEYKVFMAFVFNVV

SEQ ID	SEO ID NO:	Mer	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
l			1	sequence		
23151	53519	Α	23285	2	2556	WRKIYQAN/GKTKKAGVAILVS
						DKTDFKPTKIKRDKEGHYIMVK
						GSIQQEELTILNIYAPNTGAPRFI
1			1			KQVLSDLQRDLDSYTIIMGDFN
						TPLSTLDRSMRQKVNKNTQEL
						NSALHQADLIDIYRTLHPKSTEC
1						TFFSAPHHMYSKIDHIVGCKAL
1						LSKFKRTEIITNCLSDHSAIKLEI
1						RIKKLTONCSTTWKLNNLLLND
1						YWVHNEMKAEIKMFFETNENK
						DTTYQNLWDTFKAVCRGKFIA
	ŀ					LNAHKRKQERSKIDTLTSQLKE
			ŀ			LEKQEQTHSKASRRQEITKVRA
1						ELKEIETQKTLQKINEFRSWFFE
			ĺ			RINTIDRPLARLIKKKREKNQID
		1	İ		İ	AIKNDKGDITTDPTETQTTIREY
		l		ł		YKHLYANKLENLEEMDKFLDT
	1	l	ŀ			YTLPRLNQEEVESLNRPITGSGI
		l			1	VAIINSLPTKKSPGPDGFTAEFY
	ľ			l		QRYKEELVPFLLKLFQSIEEEGI
				l	l	LPNSFYEASIILIPKLGRDIAKKE
1			l		ŀ	NFRPISLMNIDAKILNKILAHRI
				l		QQHIKKLIHHDQVGFIPGMQG
	İ			l		WFDIRKSINVIQHINRTKDKNH
						MIISIDAEKAFDKIQQPFMLKTL
			l			NKLVLEVLVRAIRQEKEIKCIQL
	l					GKEEVKLSLFADDMIVYVENPI
						VSAQNLLKLISNFSKVSGYKIN
					ŀ	VQKSQAFLYTNNRQTESQIMSE
						LPFTIASKSIKYLGIQLTRHVKD
			ŀ	l	1	LFKDNYKPALNEIKEDTNKWK
1				1	i	NIPCSWIGRINIVKELEKTTLNFI
23152	53520	A	23286	I	3087	
23153	53521	Ā	23287	ı	741	MLGKKKPVCEVGGDSGVPKKE
		1		ľ		FOEGROVLFKLWRGOANKVG
				1		DNSIDSWKNAGRVFKDSDKFD
			İ		i	ANDPILKDQTQEWSGSATFTSD
				l		GKIRLFYTDYSGKHYGKQSLTT
	1			l		AOVKVSKS/V*HTQNORSGRSO
1	1	1	l	I		
	1			I		NDF*RRRKNIS*RSAVYR*RQL
				l		YIRRQPYAERPS\HVEDKGHKY
				l		LVFEANTGTENGYQGEESLFNK
1	l	1	l	I		AYYGGGTNFFRKESQKLQQSA
			[	l		KKRDAELANGALGIIELNNDYT
						LKKVVKPLITSN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			ŀ	sequence .		
23154	53522	Α	23288	849	1688	QQRR*MCQNLMTH\QNQPSGR
					1	SQNDF*RRRKNISERSAVYR*R
		l				QLYIRRQPYAERPS\HVEDKGH
		l				KYLVFEANTGTENGYQGEESLF
		l				NKAYYGGGTNFFRKESQKLQQ
i	1	l				SAKKRDAELANGALGIIELNND
į.		l				YTLKKVMKPLITSNTVTDEIER
		l				ANVFKMNGKWYLFTDSRGSK
		l				MTIDGINSNDIYMLGYVSNSLT
		1				GPYKPLNKTGLVLQMGLDPND
		1				VTFTYSHFAVPQAKGNNVVITS
		1				YMTNRGFFEDKKATFAPSFLM
		<u> </u>				NIKGNKTSVVKNSILEPGQLAV
23155	53523	A	23289	1	3171	MVKGSIQQEELTILNIYAPNTG
			1			APRFIKQVLSDLQRDLDSHTLI
	l	l				MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQADLIDIYRTLH PKSTEYTFFSAPHHTYSKIDHIL
1		1				GSKALLSKCKRIEITNYLSDHS
		l				AIKLELRIKNLTQSRSTTWKLN
						NLLLNDYWVHNEMKTEIKMFF
		1				ETNENKDTTYQNLWDAFKAVC
		1				RGKFIALNAYKRKEERSKIDTL
		1				TSOLKELEKOEORHSKPSRROE
		l				ITKMRAELKEIETO
23156	53524	A	23290	1	3255	
23157	53525	A	23291	1	3924	
23158	53526	Α	23292	1	3139	
23159	53527	Α	23293	1	3210	MVKGSIQQEELTILNIYAPNTG
		l				ALRFIKQVLRDLQRDLDSHTIIM
1		l				GDFHTPLSTLDRSTRQKVNKDI
						QELNSALHQEDLIDIYRTLHPKS
	1	l				TEYTFFSAPHHTYSKIDHIVGSK
						ALLSKCKRTEIITNCLSDHSAIK
					i	LELRIKNLTQNRSTTWKLNNLL
						LNDYWVHNEMKAEIKMFFETN
						ENKDTTYQNLWDTFKAVCRGK
	1	1				FIALNAHKRKQERSKIDTLTSQL
						KELEKQEQTHSKASRRQEITKIR
	ļ	L				AELKEIETQ
23160	53528	В	23294	1	3127	
23161	53529	A	23295	1	3325	
23162	53530	Α	23296	1	3145	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuelcotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amine acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nueleotide insertion)
23163	53531	A	23297	191	819	VNVTSFGSRPICSTSPVLFSGL*
						GPVKEFDT*PSM*ISFELIPSIVIF
		1		1		EPRESVNKYHLPFILKTFARSISS
						VTVFEVISGFITFFNV*SLFNSMI
				1		PRAPFANSASRFLALC*SFWLSL
				1		RKKFVPPP*YALLNKDSSPW/SS
		1				VFCSRVGFEYKVFMAFVFNVV
						RVSORMVVAGCIIAFIDKLLNV
						LICFSVSVKNRFVIFHSVDFECV
23164	53532	A	23298	I	3352	
23165	53533	A	23299	1841	2679	QQRR*MCQNLMTH\QNQRSGR
		l				SQNDF*RRRKNISERSAVYR*R
		l				QLYIRRQPYAERPS\HVEDKGH
		1				KYLVFEANTGTENGYQGEESLF
		l				NKAYYGGGTNFFRKESQKLQQ
		l				SAKKRDAELANGALGIIELNND
		l				YTLKKVMKPLITSNTVTDEIER
		1				ANVFKMNGKWYLFTDSRGSK
	İ	i				MTIDGINSNDIYMLGYVSNSLT
		1			1	GPYKPLNKTGLVLQMGLDPND
		1				VTFTYSHFAVPQAKGNNVVITS
				1		YMTNRGFFEDKKATFAPSFLM
						NIKGNKTSVVKNSILEQGQLTV
23166	53534	Α	23300	1207	1533	YVHCKPSGWTVRTFDKPRKRFI
						AFFIAGILFRAIKNHFLPRETLQ
						CLPYILTGFRRGQSEYFSIFSNM
	1	ı				DLADTVMFL*GAIRFSDLVNEQ
						IQHTFLIPGETICRLSEVV
23167	53535	Α	23301	I	3253	MGDFNTPLSTLDRSTRQKVNK
		l			1	DTQELNSALHQGDLIDIYRTLH
		l				PKSTEYTFFSAPHHTYSKIDHIV
		l				GSKALLSKWKRTEIITNYLSDH
		1				SAIKLELRIKNLTQSRSTTWKLN
		l	i			NLLLNDYWVHNEMKAEIKMFF
		l				ETNENKDTTYQNLWDAFKAVC
1		l				RGKFIALNAHKRKQERSKIDTL
	1	l				TSQLKELEKQEQTHSKASRRQE
l						ITKIRAELKEIETQKTLQKINESR
1	1	1				SWFFERINKIDRPLARLIKKKRE
			İ			KNQIDTIKNDK
23168	53536	A	23302	1	3095	
23169	53537	Α	23303	2	2678	

SEQ ID	SEQ ID NO:					Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
i	sequence		09/540,217	codon for peptide sequence	of peptide sequence	detetion, - possible nucleotide insertion)
				sequence		
23170	53538	Α	23304	1	3335	MVKGSIQQEELTILNIYAPNTG
						APRFIKQVLSDLQRDLDSHTLI
						MGDFNTPLSTLDRSTRQKVNK
1						DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQSRSTTWKLN
ľ						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
Ì						RGKFIALNVYKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
						ITKIRAELKEIETQ
23171	53539	В	23305	1	2293	
23172	53540	Α	23306	I	3810	OLIGON A LANDARING DA OR
23173	53541	Α	23307	255	1215	CVGQLAAAKR*RNSS*IQRLSR
						CVCSCGKPERR**HINLHVLSK
1						GRRQLN/QTAGKTRAVSLKTAI SSTPTIRS*KIRRKNGPVLOPLH
1						LTEKSVYSTLTIPTKAINTLYSK
						PTREOKTOTKAKNLYLTKRTT
		1			1	AAARTSSVKKARSFSRALKNA
1		1				MLS*RTAPSVS*S*IMITH*KK**
Ì		l				SR/CITSNTVTDEIERANVFKMN
						GKWYLFTDSRGSKMTIDGINSN
						DINMLGYVSNSLTGPYKPLNKT
		l				GLVLQMGLDPNDVTLTNSHFA
1		l	ĺ			VPQAKGNNVVITSYMTNRGFF
l		l				EDKKATFAPSFLMNIKGNKTSV
	1	l				VKNSILEQGQLTVN
23174	53542	A	23308	1	4539	
23175	53543	Α	23309	1	3229	
23176	53544	В	23310	1	3552	
23177	53545	A	2331I	1	3139	
23178	53546	Α	23312	I	3057	
23179	53547	Α	23313	I	3215	MGKKQNRKTGNSKTQSASPPP
		1				KERSSSPATEQSWMENDFDELR
1		1				EEGFRRSNYSELREDIQTKGKE
		1				VENFEKNLEECITRITNTEKCLK
		l				ELMELKTKARELREECRSLRSR
1				1	1	CDQLEERVSAMEDEMNEMKRE
						GKFREKRIKRNEQSLQEIWDYV
			1			KRPNLRLIGVPESDVENGTKLE
						NTLRDIIQENFPNLARQANVQIQ
1			1	1	1	EIQRTPQRYSSRRATPRHIIVRFT
		İ				KVEMKEKMLRAAREKDRSTRQ
	L					KVNKDTQELNSA

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23180	53548	Α	23314	I	1921	MVKGSIOOEELTILNTYAAHTG
						APRLIKQVLSDLQRDLDSHTIIM
				ŀ		GDFNTPLSTLDRSTRQKVNKDT
						QELKSALHQADLTDIYRTLHHK
						STEYTFFSAPHHIYSKIDHILGSK
						ALLSKCKRTEIITNYLSDHSAIK
				i		LELWIKNLTQNHSTTWELNNLI
						LNDYWVHNEMKAEIKMFFETN
						ENKDTTYHNLWDTFKAVCRGK
1		ĺ				FIPLNAHKRKQERSKIDTLTSQL
						KELEKQEQTN\HSKASRRQEITK
1						IRAELKEIETQKTL*KINESRSW
						FFERINKIDRLLARLIKKKREKN
						QIDAIKNDKGDITTDPTEIQTTIR
İ				l		EYCKHLYANKLENLEEMDKFL
				ĺ		DTYTLPRLNQEEVESLNRPITG
İ				l		AEIVAIINSLPTKKSPGPDGFTA
	1					KFYQRYKEELVPFLLKLFQSIEK
						EGILPNSFYEASIILIPKPGRDTT
i		l				KKENFRPISLMNIDAKILNKKL
1						AKRIQQHIKKLIHHDQVGFIPG
1			1			MQGWFNIRKSINVIQHINRAKD
						KNHMIISIDAEKAFDKIQQPFML
1						KTLNKL\GIKYLGIHLTRDVKDL
i						FKENYKPLLKEIKEDTNKWKNI
1		1		1		PCSWVGRINIVKMAILPKMCL*
		l				RRWTPLCLLTAAASLNRE*ERL
1						VQPLLWRQMYCGPRHCRQVPQ
		L_				HRRLNWSPSLRLS
23181	53549	B A	23315	584	2907 1278	OORR*MCONLMTH\ONORSGR
23182	33330	A	23310	384	12/8	SQNDF*RRRKNISERSAVYR*R
1		1				OLYIRROPYAERPS\HVEDKGH
i		l				KYLVFEANTGTENGYOGEESLF
1						NKAYYGGGTNFFRKESQKLQQ
						SAKKRDAELANGALGIIELNND
		1		1		YTLKKVMKPLITSNTVTDEIER
		1				ANVFKMNGKWYLFTDSRGSK
		1				MTIDVPOAKGNNVVITSYMTN
1		l				RGFFEDKKATFAPSFLMNIKGN
						KTSVVKNSILEQGQLTVN
23183	53551	A	23317	1	3345	K151 TKHSILLQOQLI YN
23183	10001	I.	23317	l <u>'                                    </u>	3343	

			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23184	53552	Α	23318	2	3366	WRKIYQANGKQKKAGVAILVS
						DKTDFKPTKIKRDKEGHYIMVK
					l	GSIQOEELTILNIYAPNTGAPTFI
						OOVLSDLORDLDSHTLIIGDFN
						TPLSTSDRSTROKVNKDTOELN
						SALHOADLIDIYRTLHPKSTEYT
	-					FFSAPHHTYSKIDHILGSKALLS
	l					KCKRTEIITNYLSDHSAIKLELM
						IKNLTQNHSTIWKLNNLLLNDY
						WVHKEMKAEIKMFFETNENKD
	İ					TT\YQNLWDTFKAVCRGKFIAL
						NAHKRK\QER
23185	53553	Α	23319	1	2611	MGDFNTPLSTLDRSTRQKANK
						DTQELNSALHQVDLIDIYRTLH
						PKSTEYTFFSAPHHTYSTTDHIL
						GSKALLSKCRRTEIITNYLSDHS
	İ					AIKLELRIKNLTQNRSTTWKLN
					1	NLLLNDYGVHNEMKAEIKMFF
						ETNENKDTTYQNLWDTFKAVC
	ĺ					RGKFIALNAHKRKQERSKIDTL
		ļ				TSQLKELEKQEQTHSKASRRQE
	İ					ITKIRAELKEIETQKTLQK\INES
						RSWFFERINKIDRPLARLIKKKR
		1				EKNQIDAIKNDKGDITTDPTEIQ
						TTIREYYKHLYANKLENLEEM
						DKFLHTYTLPRLNQEEVESLNR
						PITGSEIVAIINSLPTKKSPGPDG
						FTAEFDQRYKEELVPFLLKLFQ
						SIEKEGILPNSFYEASIILIRKPGR
						DTTKKENFRPISLMNTDAKILN
						KILAKRIQQHIKKLIHHDQVGFI
						PGMQGWFNIHKSINVIQHINRA
						KDKNHMIISIDAEKAFDKIQQPF
						MLKTLSKLGIDGTYFKIIRAIYD
		l				KPTANIILNGQKLEAFPLKTGTR
		l				QGCPLSPLLFNIVLEVLARAIRQ
				ļ		EKEIKLISNFSKVSGYKINVQKS
	1		l			QAFLYTNNRQTESQIMSQLPFTI
						ASKRIKYLGIQLTRDVKDLFKE
	1		l	l		NYKPLLKEIKEDTNKWKNIPCS
	l		l			WVGRINIMKMAILPKVIYRFNA
	1		l			IPIKLPMTFFTELEKTTFKFIWN
		l		l		QKRARITKSILSQKNKAGGITLP
	l	ı	ı	l	ı	DFKLYYKATVTKTA WYWYQN

SEQ ID	ISEO ID NO:	Mot	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	1	
		L				
23186	53554	Α	23320	1	3099	MGELITPLSTLDRSTRQKVNKD
						TQELNSALHQGDLIDIYRTLHP
				l		KSTEYTFFSAPHHTYSKIDHILG
						SKALLSKCKRTEIITNYLSDHSA
					i	IKLELRIKNLTQNRSTTWKLNN
1				l		LLLNDYWIHNEMKAEIKMFFET
				l		NENKDTTYQNLWDAFKAVCR
ŀ		İ		l		GKFIALNAHKRKQERSKIDTLT
					1	SQLKELEKQEQTHSKASRRQEI
				l		TKIRAELKEIETQKTLQKINESR
						SWFFERINKIDRPLARLIKKKRE
					ľ	KNOIDTIKNDK
23187	53555	A	23321	911	1539	VNVTSFGSRPICSTSPVLFSGL*
					ŀ	GPVKEFDT*PSM*ISFELIPSIVIF
l				ł	1	EPRESVNKYHLPFILKTFARSISS
					Ì	VTVFEVISGFITFFNV*SLFNSMI
Ī						PRAPFANSASRFLALC*SFWLSL
				l		RKKFVPPP*YALLNKDSSPW/SS
						VFCSRVGFEYKVFMAFVFNVV
						RVSORMVVAGCIJAFIDKLLNV
						LICFSVSVKNRFVIFHSVDFECV
23188	53556	Α	23322	191	820	VNVTSFGSRPICSTSPVLFSGL*
						GPVKEFDT*PSM*ISFELIPSIVIF
						EPRESVNKYHLPFILKTFARSISS
ŀ						VTVFEVISGFITFFNV*SLFNSMI
						PRAPFANSASRFLALC*SFWLSL
				1		RKKFVPPP*YALLNKDSSPW/SS
				1		VFCSRVGFEYKVFMAFVFNVV
						RVSORMVVAGCIJAFIDKLLNV
1	į.	l			i	LICFSVSVKNRFVIFHSVDFECV
23189	53557	A	23323	965	5050	TWKGTTSTSRCKIMPKYRSTRO
23,05	3333,	ľ	23323	505	5050	KVNKDTQELNSALHQADLIDIY
		l				RTLHTKSTEY/TFFSAPHHTYSK
1		l				IDHIVGSKALLSKCKRTEIITNY
1		l		l	ŀ	LSDHSAIKLELRIKNLNQSRSTT
	i			ŀ		WKLNNLLLNDYWVHNEMKAE
ŀ						IKMFFETK\ENK\DTTYQNLWD
1						AFKAVCRGKFIALNAHKRKQE
1	l		l	1		RSKIDTLTSQLKELEKQEQTHS
1	l		l	l		
1						KASRRQEITKIRAELKEIETQKT LQKINESRSWFFERINKIDRPLA
	l	ı		l		RLIKKKREENO
23190	53558	A	23324	1	2787	RLIKAKEENŲ
23190	53559	A	23324	1	902	
	53560	A	23325	1	3640	
23192	133200	IA.	23320	1	3040	

SEQ ID	SEQ ID NO:					Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23193	53561	Α	23327	1	3686	MHIVVETALSASWQNKAKPPA
						RVLLQVVPNVWFLVAVVWEL
						YPSLDLMDRSIECSSSPATEQS
		[				WTENDYDKLREEGFRRSNYSE
						LQEDIQTKGKDVENFEKNLEEC
						ITRITNTQKCLKELMELKTKAR
		1				ELREECRSLRSRCDQLEERVSV
		1				MEDEMNEMKQEGKFREKRIKR
						NEQSLQEIWDYVKRPNLRLIGV
						PESDGENGTKLKNTLQDIIQENF
		1				PNLARQAKVQIQEIQRMPQRYS
						LTRATPRHIIVRFTKVE
23194	53562	Α	23328	1	3433	MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQADLIDIYRTLH
1						PKSTEYTFFSAPHHTYSKIDHIL
						GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQNRSTTWKLN
1						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFIALNAHKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
İ						ITKIRAELKEIETQKTLQKINESR
1						SWFFERINKIDRPLARLIKKKRE
						KNQIDAIKNDK
23195	53563	Α	23329	1	3070	MEGEMNEMKREGKFREKRIKR
						NEQSLQEIWDYVKRPNLRLIGV
						PESDGENGTKLENTLQDIIQENF
		1				PNLARQANVQIQEIQRTPQRYS
						SRRATPRHIIVRFTKVEMKEKM
1						LRAAREKDRSTRQKVNKDTQE
						LNSALHQADLIDIYRTLHLKSTE
						YTFFSAPHHTYSKIDHILGSKAL
						LSKCKRTEIITNYLSDHSAIKLE
						LRIKNLTQNRSTTWKLNNLLLN
		6				DYWVHNKMKAEIKMFFETNEN
		L_				\KDTTYQNLWDA

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nuelcotide	Nucleotide location of last	Amino acid sequence ( X-Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	eodon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
		1		sequenee		
23196	53564	I .	23330	3	2372	TTNCLSDHSAIKLELRIKKLTEN
23196	33304	A	23330	]3	2312	
		l		l		HTTTWKLNNLLLNDYWVDNE
		ĺ				MKAEIKMFFEINEKK\DTTSQNI
l				l		WDTFKAVCRGKLIALNAHKRK
				l		QERSKIDTLTSQLKK\LEKQEQR
						NSKASRKQEITKIRAELKEIETQ
1					l	KKTLQKINESRSWFFEKINKIDO
ŀ						PLARLIKKKREKNQVDAIKNDK
1					l	GDIITDPTEMQTTIREYYKYLY
l				l		ANKLENLEEMDKFLNTYTLQR
		ŀ				LNQEGIESLNRPITGSEIEAIINSL
İ						PTKKSPGPDGFTAKFYQRYKEE
l		l				LVYPEPRSESECLSNIREFLRGC
		[				GASLRLESIREDRNGRSQKTVH
l						TEGDMNMNIKKIVKQATVLTF
						TTALLAGGATQAFAKENNQKA
						YKETYGVSHITRHDMLQIPKQQ
i						QNEKYQVPQFDQSTIKNIESAK
1		1				GLDVWDSWPLONADGTVAEY
1					Į.	NGYHVVFALAGSPKDADDTSI
1						YMFYQKVGDNSIDSWKNAGR
						VFKDSDKFDANDPILKDQTQE
ļ.		1				WSGSATFTSDGKIRLFYTDYSG
		1		ł		KHYGKOSLTTAOVNVSKSDDT
l		1				LKINGVEDHKTIFDGDGKTYON
		1				VOOFIDEGNYTSGDNHTLRDPH
		1				YVEDKGHKYLVFEANTGTENG
1						YQGEESLFNKAYYGGGTNFFR
		1		1		KESQKLQQSAKKRDAELANGA
1		1				LGIIELNNDYTLKKVMKPLITSN
1		1		1		TVTDEIERANVFKMNGKWYLF
l				1		
		_				TDSRGSKMTIDGINSNDIYMLG
23197	53565	A	23331	2	2987	APRFIKQVLSDLQRDLDSHTIIM
l		1		1		GDFNTPLSTLDRSTRQKVHKDT
		1		i		QELNSALHQADLIDIYRTLHPK
		1				STEYTCFSAPHHTYSKIDHIVGS
1		1			1	KALLSKCKRSEIITNCLSDHSAI
		1			i	KLELRIKKLTQNRSTTWKLNNL
		1				LLNDYWVHNEMKAEIRMFFET
		1				NENKDTTYQNLWDTFKAVCRG
		1				KFIALNAHKRKQERSKIDTLTS
		1				QLKELEKQEQTHSKASRRQEIT
		1				KIRAELKEIETQKTL\QKINE\SR
		l				SWFF\ERINK
23198	53566	Α	23332	339	701	PDKEGNEIWVDMYTVKPSGWT
[	1	1		1		VRTFDKPRKRFIAFFIAGILFRAI
		1	l	1		KNHFLPRETLOCLPYILTGFRRG
	1	l				QSEYFSIFSNMDLADTVMFL*G
		1		I		AIRFSDLVNEQIOHTFLIPGETIC
		1				RLSEVV
23199	53567	A	23333	190	502	The state of the s
23199	22307	ΙΛ.	23333	170	302	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	ł	
23200	53568	Α	23334	4472	8032	KKSGNNRLMRVHGKKRFLWQ
						KQGVCEVEIQAQRESARDGTD
		1		ŀ		REDAESPHAEVTVPRGTRTEILT
				l		LWASRDRPASACGVTVPKGST
				l		THPGGMDRVCGVDTAVLHWM
				l		KQVQGISNLEVETGPRRRDMD
						EAGSRHSQQTNIGTENQTPHVL
				l		THKWQLNRNSACDKLHKPHSV
		l				PGGVAAGDVGGKTISNPGVCAI
						EYQVDGSPHIWEDVNKGVYIEE
				l		DRAPLMERSSSPAMEQSWMEN
						DFDELREEGFRRSNYSELRED
23201	53569	Α	23335	1	2374	MVSISDLVICPPRHPKVLGLQGP
						PGLDSISDPSAGAGFLDWGEIG
	ŀ			l		MPGPGRAGHQALCKCDCQCLE
		l				KTTTKAPGKMPKSTRSGPVRV
		l		l		RLADGPNRCAGRLECGMPDAG
				l		EQCVMTTGTSGRHCGLLGTGL
						WKGYTDLTIIPPGPGTPPQERTC
		1				QGDYHSGGTWTHSPLETTRRP
		l			{	GSSSPAIRRLPAQMLLLPARPPH
	i					PRSSSPEAMDPPPPKAPPFPKAE
			l		1	GPSSTPSSAAGPRPPRLGRHLLI
		1				DAN/GVYPYTYTVQLEEEPRGP
						PQREAPPGEPGPRKGYSCPECA
					l	RVFASPLRLQSHRVSHSDLKPF
						TCGACGKAFKRSSHLSRHRATH
						RARAGPPHTCPLCPRRFQDAAE
						LAQHSWGTPRGPLLAAACNCE
						VARGRLESPGPERLLHGYGGRE
						EEGGWGRAAGGLDRVEGFISS
						KAHHYLLIDTQGVPYTVLVTRS
		1		l		HRGSQGPVGLQARKVLQLPRV
					1	LKGLRVHVHLQRHSITHSEVPQ
	1					DFAGSLDSFQTPGESLRLVFRA
						LDTTQSSRISKAEPCLKEEPLSL
				i		GDLPYMHTTLCFCRKRRASPGP
					1	GTLQRGALAWPDWASPRALPV
	1			I		PSLSSTTRSPAAPLFAVPLSGRT
	l		1			TQAMAFDGIIFQGQSQRSAGLT
	1			I		TTSRFLACQRPLRLCAWWASRS
				I	1	PRCTLRRPVGLRPGVHPRPRLV
		1		l		YRDLKPENVMASGOPRDRPOP
	I	l	1	I		WFAWPPRPTRFCGGCWTLTPK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codnn, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23202	53570	A	23336	853	1602	PGAAWSRPDLRGCCTEPQPALR
23202	33370	<u> </u> ^_	25550	055	1002	MLVLASPCPOPLAFSSVETMEG
i		ı				PPRRTCRSPEPGPSSSIGSPOASS
						PPRPNHYLLIDTQGVP\YTVAW
1						LDEESOREPGASGAPGOKKCYS
						CPVCSRVFEYMSYLQRH\RFSH
1						SEVKPFE\CDICGK\AFKRSQHL/
1						SARHHSIHLGGWVGGPTACRFC
						PRRF\RDAGELAQHSRVHSGER
	İ					PVSVSTLPSPLYGAEHTAETHA
						VEASHEAGLPGAPGTHRTLQG
		1				ALDSCPDTW
23203	53571	A	23337	1	1144	MGHHLMIDKLVALGGLYYAIQ
						RHYATKCSVLKNDQILVIGLFM
1						IQNVIYRKHFANPLSALFLQQGI
						ELFAAIAEIHITTVAERNHAITQI
						RLEAQTIFDSLKEWHNAIRKSL
						DYEALQAFEFRVGATDRGFPAL
						SSEALVRVLVLDANDNSPFVLY
		l				PLQNGSAPCTELVPRAAEPGYL
		1				VTKVVAVDGDSGQNAWLSYQ
		ŀ				LLKATEPGLFGVWAHNGEVRT
					l	ARLLSERDVAKHRLVVLVKDN
						GEPPRSATATLQVLLVDGFSQP
						YLPLPRAAPAQAQADSLTVYL
						VVALASVSSLFLFSVLLFVAVR\
						LCRRSRAAPVGRCSVPECPFPG
	l	l				HLVDVSGTGTLSQSYQYEVCLT
						GGSGANEFKFLKPVIPNLLSRDS
23204	53572	Α	23338	1	1260	MVCYIQENLPFLLKPSVENFYIL
						ITEGALDREIRAEYNITITVTDL
						GTPRLKTEHNITVLVSDVNDNA
		1				PAFTQTSYTLFVRENNSPALHIG
					1	SVSATDRDSGTNAQVTYSLLPP
1						QDPHLPLASLVSINADNGHLFA
						LRSLDYEALQAFEFRVGATDRG
1		1				SPALSREALVRVLVLDANDNL\
						PFVLYPLQNGSAPCTELVPRAA
		1				EPGYLVTKVVA VDGDSGQNA
		1				WLSYQLLKATEPGLFGVWAHN
1			1	1	l	GEVRTARLLSERDAAKHRLVV
	1	1	1		l	LVKDNGEPPRSATATLHVLLVD
1						GFSQPYLPLPEAAPAQAQAEAD
			1	1		LLTVYLVVALASVSSLFLLSVL
		1		1		LFVAVRLC\RRSRAASVGRCSM
			1	1		PEGPFPGRLVD\VSGTGTLFQSY
						QYEVCLTGGSETGEFKFLKPITP
	L		L	l		HLPPHRGGKEIEENSTLPNSFGF

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X-Unknown, *=Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23205	53573	A	23339	123	5794	SSSDSGSLKGACTEVKMEPAGE RFPEQRQVLILLLLLEVTLAGW EPRRYSVMEETERGSFVANILAN DLGLGVGELAERGARVVSEDN EQGLQLDLQTGQLLINEKLDRE KLGGFTEPCIMIFQVLLKKPLE VFRAELLVTDINDHSPEFPERE MTLKIPETSSLGTVFPLKKARD LDVGSNNVQNYNISPNSHFHVS TRTRGDGRKYPELVLDTELDRE EQAELRILTIAVDGGSPRSGT VOILLIVLANDNA
23206	53574	C	23340	20	313	
23207	53575	Α	23341	119	247	RSRCISASPRRPASFRKSIPEEED ESLNI*KGQVSILFFYIK
23208	53576	В	23342	55	288	
23209	53577	A	23343	1	3576	MAFLFEFSDLVSLLSIGTMLAY SLVAFSVLVLRYQPDENFSKNE KPKEEVVBMNSVPKAESPACVP EASSTPASLWSPVSTIPTPRWGR IVYGCAFILVVLLSMLCLVLAH WPKRLFSGELIVIAAAVLLLVLI VGFTFTVWRQPQSNTFLVFKEP HISTSQNLSLPEHPMEYLLRTGI EPDVRGVLTVNALPAGPPVACS PLVSISAHVYLMMQMTTETWA QCGVWMLIALECPNLKLKKPP WLIMPLAMTMYA

3059

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X-Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23210	53578	A	23344	3	1762	SVREOFGHIVRVRFFSI.SPLPLR PWPULCPAR.RTMGSILSRRIAG VEDIDIQANSAYRYPPKSGNYF ASHFFMGGEKFDTPHEGHLIFG ENMDLNFILGSRPSQFPYVTPAP HEFVETTLRSI,VNIRKDSLRLVR YKDDADSPTEDGDKPRVLYSL EFTFDADARVAITIYCQASEEFL NGRAVYSFKSPSLQSETVHYKR GYSQQVSLPSFKIDFSSWKDDE LNFDLDRGVFPVVIQAVVDEG/ DSVVEVTGHAHVLLAGFEKHM DGSFSVYRLEXDGKOVDRVSYLL QEIYGIENKNNQETKPSDDENS DNSHECVVCLSDLRDTLLFC HLCLCTSCADTLRYQANNCPIC RLPFRALLQIRAVRKFOALSP YSSPVLAGSLEHDEHSGPFKK SKPHFASLASKKPKRETNSDSV PPGYEPISLLEALNGLRAVSPAI PSAPLYEETTYSGISDGLSQASC PLAAIDHILDSSRQKGRPGSKAP PSAPLYEETTYSGISDGLSQASC PLAAIDHILDSSRQKGRPGSKAP DSTLRSYSSPHEEDEEKIS.EDV DAPPPLGGAELALRESSPESFI TEEVDESSSPQGTRAASIENVL QUSSSPEHCGGGPPADIVLPALG
23211	53579	A	23345	35	3134	RPREGPTWPQLPGSSPRRLEPW SPGSSSRGLFPWSPGSSSRGLF WSPGSSSRGLFPWSPGSSSRGLF PWSPGSSSRGLFPWSPGSSSRG GSSRGHPGPPPGGSSRGF GSSRGHPGPPPGGSSRGHPGPP GALPVVTRVLLPGLFPWSPGSSSRGHPGPPWSPGSSSRGLFPWSPGSSRGHPGPPWSPGSSRGHPWSPGSSRGHPWSPGSSRGHPWSPGSSRGHPWSPGSSRGHPWSPGSSRGHPWSPGSSRGHPWSPGSSRGHPWSPGALPVVTRVLLBGTPWSPGSSRGHPWSPGALPVVTRVLLBGTLPVVTRVLLBGTLPVVTRVLL
23212	53580	A	23346	3	91	7.11.122
23213	53581	A	23347	3	174	
23214	53582	A	23348	3	360	
23215	53583	В	23349	158	368	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		***
23216	53584	A	23350	3786	4928	PPPARTRPCASLTPEKAKWWRS
		1				KPGLTRAPARCGLSSPOTGSCS
	1	1		1		APASAG*VSGNSRSGTRRGLRP
		l				TRG*GPCGPSSRARAISSPRAS/L
		l				RMSQRELGLWDPNNFEEPVAL
						QEMDTSNGVLLPFYDPDSSIVY
		1				LCGKVLTAGQGEQGTGWRCG
		1				GPCPGAPLNRLILQGDSSIRYFE
		1				TDEPPFVHYLNTFSSKEPORGM
		1				GFMPKRGLDVSKCEIARFYKLH
		l		i	Ì	ERKCEPHMTVPRKSDLFQDDL
						YPDTPGPEPALEADEWLSGQD
					İ	AEPVLISLRDGYVPPKHRELRV
						TKRNILDVRPPSGPRRSQSASDA
						PLSVRSALLHSGPIYISPNRPYTT
				l		RPLCPPPQQQHTLETLLEEIKAL
						RERVQAQEQRITALENMLCELV
23217	53585	Α	23351	1	1898	MSRKGPRAEVCADCSAPVGPD
						FALAISELVSPDIFMQSHSENAIS
				ì		VKEIVTEIESISQGVGQIQLKGDI
i						LPNPCHTPKKNSIHELLLERAQT
l		i				PENKPGHMEQDEDSCTAQPEL
						AKDSGMCNPEGCLTTHSSIADL
				ľ		EEGEQLRGNKSSRAQGCTQNS
İ				į.		KNDSSVADLAPKGKSDEAPPEH
		1		1	1	SFVLKEPEMSKGKGKYSGSEA
1		1				GSLSHSEQNATVPAPRVLEFDH
						LPDPQEGPGPEASNGLKVSQKS
1						EGSQPSSSRSPLEFLKEAGVPNQ
ł		1				TLSSGTPDPYFMALLGPKLNSR
						RPHLHPVSHTLDLGNSPPLPQQ
						FAKCQRGRKGPEGKYPLHYLV
				i		WHNRHRELEKEVRAGQTLITQ
		1				TLSQANPTAITAEEYFNPNFELC
						NRDMGRPMELTTKTQKFKAKL
		1		i		WLCEEHPLSLCEQVAPIIDLMA
l		1		1		VSNALFAKLRDFITLQIPIFHILN
1		1		1	1	ARITFGNLNGCDEPVPSVRGSPS
ļ.		1		1	}	SETPSPGSDSSSVSSSSSTSEAPR
	1		1			ENACPSALPGVASCRGCEISPAL
						FEAPRGYSMMGGQREAATRDD
1						DDDLLQFAIQQSLLEAGSEYDQ
1	1	1	1		[	VTIWEALTNSKPGTHPMSYE/R
1					1	SPTGQGQVPAATCSG/RYDEQL
1		1				RLAMELSAQEQEERRRRARQE
		١	00000			EEELERILRLSLTEQ
23218	53586	Α	23352	341	711	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ł		sequence		
		ᆫ				L
23219	53587	Α	23353	57	719	TSDLMAEIIQERIEDRLPELEQL
		1				ERIGLFSHAEIKAIIKKASDLEY
		l	1			KIQRRTLFKEDFINYVQRIGYSF
		1				KKDEIENSIVHRVQGVFQRASA
		1				KWKDDVQLWLSYVAFCKKWA
		1				TKTRLSKVFSAMLAIHSNKPAL
i		l				WIMAAKWEMEDRLSSESARQL
i		1				FL\RALRFHPECPKLYKEYFRM
		1				ELMHAGKTGGRRRKNLKKPV
		l				WMWRILIILKKSLRASWHGSST
23220	53588	A	23354	27	352	TESVVTIMSLLNKPKSEMTPEE\
23220	33300	ľ	25554	["	1552	REEEFNTGPLSVLTQSVKNNT
		l				OVLINCRNNKKLLGRVKAFDR
		l				HCNMVLE\NVKEMLTEG/DRYI
		l				
23221	53589	A	23355	32	575	SKMFL\AGDPVIVVLRNPLIAGK
23221	133389	A	23333	32	373	STRLDLPKCWDYRSLFNKP\KS
		1				EMTP\EELQKREE\EEFNT\GPTP
	l	l				LCLTQS\VKNNTQSAQSTC\RNN
		l				KKLPGAA*RPFDRASPPCSPPGT
Į.	ŀ	l				CNMVL\ENVK\EMWT*GTQEV
	ŀ	l				GKGKEGSSKP\VNKDRLTSSKM
		l				FPGGGGFSSFVVPCGNPFHSAG
		l				QIRGPPVLFDKNSHPLVLLKTA
						CHLVFENY
23222	53590	Α	23356	103	713	VTSGCGKKKVWLDPNET\NEIA
i	l	l		1		NANFRSADPGRLIKDGLIIRKPV
				1		T\VHSRARLPEKTPLARCKGHG
						TLGIGKRKGTANA\RMPKEGHH
		l				GMRRMRILAPACSRRYRESKKI
						D\RHMYHSLYLKVKG\NVFKNK
	l				1	ADFSWEHIHKAEGRARPRKEAS
		1	ŀ		1	WLDQAEA\RRSKDQGKHGKRR
	i					EERLPGORKEEINORLYSKEGR
						RPRNKNLTL
23223	53591	Α	23357	1	2847	
23224	53592	Α	23358	I	2823	
23225	53593	Α	23359	3	2858	AMFGFQRRGLGTPRLQLWLLL
		1				LEFWEVGSGOLHYSVSEEAKH
1	l	l	1		1	GTFVGRIAQDLGLELAELVQRL
1	1	1				FRVASKTHGDLLEVNLONGILF
1		l				VNSRIDREELCGQSAECSIHLEV
1		l	1			IVDRPLOVFHVNVEVKDINDNP
1		l				PVFSLREQKLLIAESKQSDSRFP
1	l	l		l		LEGASDADIEENALLTYRLSKN
1	1	1		I	i	
		1				EYFSLDSPTNGKQIKRLSLILKK
	1	l	1		l	SLDREKTPELNLLLTATDGGKP
1		I		1		ELTGTVRLLVQVLDVNDNDPE
L	L	L_		l		FDKSEYKVSLMEN

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23226	53594	A	23360	194	931	NSPVVPVMASGQFVNKLQEEVI CPICLDILOKPVTIDCGHNFCPO
						CITQIGETSCGFFKCPLCKTSVR\
					ŀ	RDAIRFNSAVAEIWWEKIQALQ
					İ	ASEVQSKRK\EAT\CPRHQEMF
						HYFCEDDGKFLCFVCRESKDH
						KSHNVSLIEEAAQNYQGQIQEQ
						IQVLQQKEKETVQVKAQGVHR
						VDVFTDQVEHEKQRILTEFELL
						HQVLEEEKNFLLSRIYWLGHEG
						TEAGKHYVEIPLMPTVERSQEA
23227	53595	Α	23361	2	381	AINWTSARPVSQCWDP\CPPSQ
						VVCLAGLPP*RVCQNPVKGP\N
						CPLAHTSEPTGHCLL*TRGMCF
						PSLCGQLCP*SGPPDCPSPRALH
						HTGFCPVALLVAESQVRARAC
						GPEPHRAGRASRWCLHAG
23228	53596	Α	23362	213	574	PQTHTGNHSERGRQPITRSPEKP
						EP\CPSKRAPQTQQP\PWAPSLC
						PPALGPCKGTRRPRVQVHRPTS
						SQKCPLPAWFHHLPLWPTSAAR
						SSSTETPTGQQMRKLRHPAAPR
						SANALSLGE_
23229	53597	Α	23363	21	456	
23230	53598	Α	23364	2	98	
23231	53599	Α	23365	64	241	PWPLASNPRGCLP*PSTLN*PTP
		l	l	l		SSQPGMSLDSLDRHPP*TRETAS
						GIGGQRPGSDPARTPHFISQTQ
23232	53600	Α	23366	209	353	

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SEQ ID	Toro In NO.	124.4	LOCO ID NO.	Ter to all do	In	I
NO:	of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence	1.00	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23233	53601	Α	23367	1766	3292	SSCPTTRVLIMKGAGHPCYLDK
		l				PEEWHTGLLDFLQGLQLWLMH
		l	l			MOQLMSLAVVSGRLSRGQCGT
		l				ERPSRRAAAAAAAGOPORGHF
	1	1				OHIEERLETNVONCOSPDTLCP
	İ	l				THCSPVLNEPCPEVSVVAVAPA
		ĺ				SPDRMSGSDGGLEEEPELSITLT
		ļ				LRMLMHGKEVGSIIGKKGETV
		1				KRIREOSSARITISEGSCPERITTI
		1				TGSTAAVFHAVSMIAFKLDEDL
		1				CAAPANGGNVSRPPVTLRLVIP
		1				ASOCGSLIGKAGTKIKEIRETTG
		1				AQVQVAGDLLPNSTERAVTVS
		1				GVPDAIILCVRQICAVILEGFSV
		1				QGQYGAVTPAEVTKLQQLSSH
		1	1	ı		AVPFATPSVVPGLDPGTOTSSO
		1	}			EFLVPNDVSRGWDGWRLIGCVI
		1				GRQG\TKINEIRQMSGAHIKIGN
		ı				O\AEGAGERHV\TITGSPVSIALA
		l				QYLITACLETAKSTSGGTPSSAP
		l				ADLPAPFSPPLTALPTASPLAC
		l				WAHPMPSPSPTLIGFKAHALLG
						FTTCFPRAAAGLGGLHRODGSS
23234	53602	В	23368	1	363	
23235	53603	A	23369	486	947	LDGAGLLAPVKAHLOKKGPRL
		1				GTRGCAMPRLGSTPFLKGQCL
		l			1	ANSRNTLEAVSLTESSASRVPP
		l				APVLGSRRPHSPSPWP/P/YYTP
						VPPPPRMALVPPNMHGQGTVG
						LTSVPLACTVAMLGSYKQAGA
						FGPLANCYGGLGHAQCWGQEF
		1				СРТОСР
23236	53604	A	23370	38	680	GSRLRRLOAAAARPALPLPLPP
						WEWKHLPHVPEAKWWLTTAR
1		t				HSAAYRADPLRVSSRDKLTEM
1		İ				AASSOGNFEGNFESLDLAEFAK
ĺ			ŀ			KQPWWRKLFGQESGPSAEKYS
						VATOLFIGGVTGWCTGFIFOKV
						GKLAATAVGGGFFLLQLANHT
						GYIKVDWQRVEKDMKKAKEQ
		l	1			LKIRKSNQIPTEVRSKAEEVVSF
						VKKNVLVT\GGFFGGFLLGMAS
23237	53605	С	23371	30	200	
23238	53606	Ā	23372	14	161	
23230	122000	<u>'``</u>	20012		101	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	ĺ			sequence		
23239	53607	Α	23373	69	715	APALPGCEHMMAIRELKVCLL
		l		1		GDTGVGKSSIVWRFVQDHFDH
		1				NISPTIG\ASFMTKTVP\WENDFT
		l				KFLIWDTAGQERFSFIGLPLYYS
		l				RLQLAAVYSCYGYLPSRDFIFIT
		l				LKE\WV\KRLKELGPE\NIVMAI
		1				AGNKCDLSDI\REVPPEGMLKE
		1				Y\AESIGAIVVETSAKNAINI\EE
		1	į		1	LFQGIS\RQIPPLDPPMKNGKQW
						DHOKLRSOPMOASRRCC
23240	53608	A	23374	3	293	VTSTMAYRAQGQKV\QKVMV
						QPINLIFRYLQNRSRIQVWLYEQ
l		1				VNMRIEGCIIGFDEYMNLVLDD
		1		1		AEEIHSKTKSRKOLGRIMLKGD
		1				NITLLQSVSN
23241	53609	A	23375	3	393	HESCSQRQRAGCALCENSHPW
		ļ.,		ľ		RSRGPGVQKSAEGLWCOPICRT
		l	İ			SSSRYL\QNRIAGFQVWLYEQV
l		l				NM\RI\EGCIIGF\DEYVNLVLDD
		ì				AEEIHSKTKSRK\OLGR\IMLK\G
		1				DNITLLQTCLPPRMVHGSGEIV
23242	53610	С	23376	129	425	
23243	53611	Α	23377	470	635	ENSSSHHIIKPFSSGCSIHGFSGH
		1				INVTSVTIL*PIKSCHIRCHF*CA
						QYMVADN
23244	53612	Α	23378	7I	566	PLFPPCLSPVVAGWARGHMLG
				1		QSHQQVAAPLKHRPITPTELCQ
		1				KCLSLIKKTKFLNAYITVSEEVA
						LKHAEESEKRYKNGQSLGDLD
l		1				GIPIAEKDNFSTSGIETT\CASNM
		ı				LKGYIPPYNATVDQKLLDQGA
ĺ						LLMGKTKLNDFAMGSGSTDGV
						FGPVKNPWSYSK
23245	53613	Α	23379	255	1042	ILAARMGKQNSKLRPEVMQDL
1	l	1	ļ	l		LESTDFTEHEIQEWYKGFLRDC
	l	1		l	1	PSGHLSMEEFKKIYGNFFPYG\D
1		1	l	1		ASKFAEH\VF\RTFDANGDG\TID
	1	1	l	l	l	FREFIIALSVTSRGKLEQKLKW
	l	1		1	i .	AFSMYDLDGNGYISKAEMLEIV
ŀ	ŀ	1				OAIYKMVSSVMKMPEDESTPE
1	l .	1	l		l	KRTEKIFROMDTNRDGTWVPH
ŀ		1	l			LALIYRTAEGGFAVORASVSFY
1		1	l	l		PPPSAGNAMVNEAAVFLEIPLG
1		1	l			SSPGOVSAVCHSGRKLSLEEFIR
1		1	l			GAKSDPSIVRLLQCDPSSAGQF
	l			L	L	G. H.C.D. G. T. C.D.QCDI GG/TOQ1

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
23246	53614	A	23380	1	721	MDGNLRKIRFEKHAITEEFRTT
23240	33014	^	23300	l'	1/21	EPGLAALRLPSTWIRGVRTSES
	İ	l				KREVGKGGOSOOYENRDTEGE
	l	l				GHVMTKARDWSKAAASQRLP
	1	ŀ			i	
						ANHLKLRLCIDGFRSHSLPEEGP
1		l				KKPCQHHAAQEEPCTRSGRLPQ
		l				AHEGSISDCCTSSEQGSMDIGSV
1		1			İ	EPGTGYNLLVCHLL/STLEKRSI
		1				WAGVSRFSRYHLSRLPLARK\G
		1				NPLTPCASQLLLPLPSTTQKRQS
						GHALYSFPYRPQPKNDNLVMP
23247	53615	Α	23381	80	562	DRAMWGDLWLLPPASANPGT
	1					GTEAEFEKAAEEVRHLKT/RSPS
		1				DEEMLFIYGHYKQATVGDINTE
1	i .					RPGMLDF\TGKA\KWD\AWNEL
						KGTSKGDAVEA\YLNQV\EELK
	İ			i	ŀ	KKYGDMERLDLVTVPCVYPKP
	i					ETIPFFFSNTVDGWEFRENNPV
						KPATQRLLTHTGS
23248	53616	Α	23382	380	1041	EVTINSGFIFIMLFMEHPEHYTI
		l				MHKWILTWILPTLLYRSCFHIIC
	i					LVGTISLACND\MTPEQMATNV
İ		1				NCSSPERHTRSYDYMEGGDIRV
	l					RRLFCRTQWYLRIDKRGKVKG
	l	1				TQEMKNNYNIMEIRTVAVGIVA
		l			i	IKGVESEF\YLAMNKEGKL\YA
	ŀ	l			1	KKECN\ED\CNFKELILENHYNT
		l				YASAKL/WTHNGGEMFVALNO
		l				KGIPVRGKKTKKEOKTAHFLP
		l				MAIT
23249	53617	Α	23383	127	549	SSPKMAATMKKAAAEDVNVTF
	1	1				EDQ\QKINKFARNTSRITELKEEI
		l				EVKK\KQLQNL\EDACDDIMLA
	1					DDD\CL\MIPY\OLGDVFISHSOE
		1				ETQEMLEEAKKNLQEEIDALES
						RVESIORVLADLKVOLYAKF\G
						SNINLEADES
		1_				SNINLEADES

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SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23250	53618	Α	23384	1	2050	MEYESMVISFDDRVMEKTEYG
		1				PIFTVFAMGNRMTFVTEEEGIN
				•		VFLKSKKVDFELAVQNIVYRT/
						SIPKNVFLALHEKLYIMLKGKM
						GTVNLHQFTGQLTEELHEQLEN
						LGTHGTMDLNNLVRHLLYPVT
						VNMLFNKSLFSTNKKKIKEFHQ
		1				YFQVYDEDFEYGSQLPECLLRN
						WSKSKKWFLELFEKNIPDIKAC
					1	KSAKDNSMDSDEDPVNLQSRS
						ALVCEMLHVQRSSSDADAIGPI
						ADHSLSNRWEVILADEHGIIAD
						DMISRYALSEKSQVELSTQVIKS
		1				ASSKSVNKSDIDTSVFLNWYNF
	1				EKKMIRAYATVIDGPEYFWCQI	
	ı				ADTEKLQCLEVEVQTAGEQVA	
	1	1			ł	DRRNCIPCPYIGDPCIVRYREDG
		1				HYYRALITNICEDYLVSVRLVD
		ı				FGNIEDCVDPKALWAIPSELLS
		1				VPMQAFPCCLSGFNISEGLCSQ
		1				EGNDYFYEIITEDVLEITILEIRR
	1				DVCDIPLAIVDLKSKGKSINEK	
	1				MEKYSKTGIKSALPYENIDSEIK	
	1				QTLGSYNLDVGLKKLSNKAVQ	
l		1		i		NKIYMEQQTDELAEITEKDVNI
	!			1		GTKPSNFRDPKTDNICEGFENP
l	ļ	1				CKDKIDTEELEGVSQKAQESMO
l						TEDMRKSSCVESFDDQRRMSL
l		1				HLHGADCDPKTQNEMNICEEE
		ł	1			VEYKNRDAISALMP\FSLRKKA VMEASTIMVYQIISQLNYRTPTI
23251	53619	A	23385	3	602	TRPPTRPGLSPDPTSPTEADRGV
23231	33019	^	23363	ľ	002	CERPHRCFLLPQITHPSSGRRCL
l						CLLSPTSFDFSLLYESOSDRAIF
		1				GVGTEPDQEPPPPYHEQVPVSV
		1				YHPT\P\SQTWVATQLTEEEQIR
		1				AQRIGLIQHLPKGVY\DPWRDG
		1	١.			SEKKIRECVICMMDFVYGDPIR
				_		FLPCVHEWYHLDCIDDWLMRS
		1				FTCPSCMEPVDAALLSSYETN
23252	53620	la-	23386	33	512	TAARCQGGPTHVPAFVGTACP
23232	153020	1	23300	155	l	TRIPGHAPAROCREYYY*YYYE
		1		1		F/WKEPRRCHALLLPPWAGLEV
		1			1	GMPPSHTQLGMHPAPPPPHLEV
		1		1	1	KAQG\PSNRQRTDGCRWCSGR
		1				ASALGGGRODSRLRERGGGYD
		1		1	1	H/HCPICRSLKIOLKCHPASGPO
		1		1	l	TVGLGRLASVS
23253	53621	A	23387	1	1731	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:		Nucleotide location of last	Amino acid sequence ( X-Unknown,
NO:	of peptide	hod	in USSN	tocation of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
23255	53623	Α	23389	ī	1986	MGIGEGPTTSEGCRESVEYSHQ
				ŀ		KSELPLSGDSQVPGPIGCNCDP
						QGSVSSQCDAAGQCQCKAQVE
						GLTCSHCRPHHFHLSASNPDGC
						LPCFCMGITQQCASSAYTRHLIS
						THFAP/GDFQ/GFA/LVNPQRNS
						RLTGEFTVEPVPEGAOLSFGNF
						AOLGHESFYWOLPETYOGDKG
						EAYFARMRRAHONRTSLSEEO
				1		LRAAVTAGRIPEPPEGRDWAQR
						ASOFSLSYEGFSLLPGSLYYWO
						LPRAFLGDKVAAYGGKLRYTL
						SYTAGPOGSPLSDPDVQITGNNI
						MLVASQPALQGPERRSYEIMFR
i			İ	l		EEFWRRPDGQPATREHLLMAL
1	ŀ		i	l		ADLDELLIRATESSVPLAASISA
ĺ		l		i		VSLEVAQPGPSNRPRALEVEEC
i		1		i		RCPPGYIGLSCODCAPGYTRTG
ł	ŀ	l		ĺ		SGLYLGHCELCECNGHSDLCHP
1						ETGACSQCQHNAAGEFCELCA
		l				PGYYGDATAGTPEDCOPCACP
		l				LTNPENMFSRTCESLGAGGYRC
		ı	ļ			TACEPGYTGQYCEQCGPGYVG
ŀ		l				NPSVQGGQCLPESSELHFPSVQ
ŀ		l				PSDAGVYICTCRNLHOSNTSRA
		Į			1	ELLVTEAPSKPITVTVEEQRSQS
					I	VRPGADVTFICTAKSKSPAYTL
		l			i	VWTRLHNGKLPTRAMDFNGIL
						TIRNVQLSDAGTYVCTGSNMF
1						AMDOGTATLHVQGSAEAGERS
		l				MYSISVLLKYTS
23256	53624	A	23390	3	13319	PASGRIRAARSGRRGASERESG
23230	33024	ľ`	23370	Ĭ	13317	AGRAMGWRAPGALLLALLLH
1						GRLLAVTHGLRAYDGLSLPEDI
1		1			1	ETVTASQMRWTHSYLSDDEYM
		1			1	LADSISGDDLGSGDLGSGDFQM
					1	VYFRALVNFTRSIEYSPOLEDA
ŀ					İ	GSREFREVSEAVVDTLESEYLKI
						PGDQVVSVVFIKELDGWVFVE
1						LDVGSEGNADGAOIOEMLLRVI
	1		1			SSGSVASYVTSPQGFQFRRLGT
	1		1			VPOFPRACTEAEFACHSYNECV
	1	1	1		1	ALEYRCDRRPDCRDMS
	<u> </u>		L	L		MLE I KUDKKPDUKDIVIS

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23257	53625	A	23391	41	1323	HYLVFFLPSLRRYTVSLSPNSTS YQLLVPGPSY/SALPISSAASLY AGA/GGGGSSRMSVSHSTSLRDG ISSGGLAVTGMARGLGGMGGIQ NKKETTQSLNDSLAVRVRSLET ENRKLESKIREHLEKKGPQVRD WSHYFKTIEDLRAQIFANTVDN VRIFQIDNARLAADDFRVKYE TELAMRQSVESDIVGLHKVIDD TNVINGCSFFMKKNHGEEVKC LQAQIASSGLTMEVDAPKSQDL AKTTADISTRYDELARKNREEL DKYWSQPIEESTTEVTMQSAKV RAAEMTFMELRRTVQSLEIDLD LMRNPKASLENSLREVEACYA LQMEQL/NLDP/GILLHLESELTQ
						LEAEIATYCLLLEDGKDFNLGD TLDSSNPMQTIQKTTTSRIVDG KVVSETNDTKVWRH
23258	53626	Α	23392	1	77 I	
23259	53627	A	23393	732	1656	SCSWTSPAAWPPNASLHSCLKR RNOPHLLWUWRFLDFSSMA SKCQSPLLPQKKKPRPLPALGP EKTSAAGLPKKGEKEQQETIE HIHEVQNEMDKINEQAKRSEL ISKIPDFWYTFVNHPEVSALPG EEDEEALHYLTRVEMTDFDDIK SGYRIDFYFDENPYFENNILAKE FHLNESGDPSSKSTEIKWKYGK DLMKRSSGOPINKAHRKRQHEE PESFTWFTDHSDVGADELGEV IKDIWPNLQYVLVPDMONEGE GEGEDDDDDEEEGGLEDIDEEG DEDGGEGEDEGERGEEDBGE DEDGGEGEDBGERGEEDBGE DEDGGEGEDBGERGEEDBGE
23260	53628	A	23394	3	923	ARRLYWWFWDFPNSMAPKRGS PLPPOKKKPRPPPALGBEETS AGLPKKGEKEQGAIEHIDEVQ NEIDRLNEQASEELIKVECKYN KLRQPFFCKRSKLISAGIPNFWV TTFVNIHPQVS\ALLGEEDEEAL\ HYLITRVEVTEFEDIKSGYRIDF FYDENPYFSKVLISKEFILINES GDPSSKSTEIKWKSGKDLTKRS SQTQNKASRKRQHEEPESFFTW FTDHSDAGADELGEVIKDDIWP NPLQYYLVPDMDDEEGGEED DDDDEEEGLEDIDEEGDEED EDEDDDGEEEGLEDIDEEGDEED

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	bod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23261	53629	Α	23395	732	1656	SCSWTSPAA WPPNASLHSCLKI
				1		RNQDHLLLWDWRFLDFPSSMA
	ŀ		1	1		SKCOSPLLPOKKKPRPLPALGP
	l					EKTSASAGLPKKGEKEOOETIE
				1		HIHEVONEMDRLNEOAK\RSEL
				İ		ISKIPDFWVTIFVNHPEVSALPG
				l		EEDEEALHYLTRVEMTDFDDIK
	İ	ĺ		l		SGYRIDFYFDENPYFENNILAKI
						FHLNESGDPSSKSTEIKWKYGK
						DLMKRSSQTQNKAHRKRQHE
						PESFFTWFTDHSDVGADELGEV
				1		IKDIWPNPLQYYLVPDMDNEE
		1				EGEEDDDDDEEEEGLEDIDEEG
		1				DEDEGEEDEDEGEKGEEDEGE
23262	53630	A	23396	3	924	ARRLVVWFWDFPNSMAPKROS
23202	33030	<u> </u> ^	23390	ľ	724	PLPPQKKKPRPPPALGPEETSAS
						AGLPKKGEKEQQ\EAIEHIDEVO
l						NEIDRLNEQASEEILKVEQKYN
		1				KL\RQPFFQKRSK\LSAGIPNFW
					1	VTTFVN\HPQVS\ALLGEEDEEA
ļ						L\HYL\TRVEVTEFEDIKSGYRID
ļ						FYFDENPYFENKVLSKEFHLNE
						SGDPSSKSTEIKWKSGKDLTKR
ļ		1				SSQTQNKASRKRQHEEPESFFT
		l			i e	WFTDHSDAGADELGEVIKDDI
ŀ						WPNPLQYYLVPDMDDEEGEGE
	1		}			EDDDDDEEEEGLEDIDEEGDED
						EGEEDEDDDEGEEGEEDEGED
23263	53631	A	23397	1	1393	MGKQKGGGPQAWEAEGLGRE
23203	33031	^	23351	l'	1555	WMRMPQEGKGSQRKSTQKQT
1				l		KREIDA YIVQAKERSYETVLSF
						GKRGLNIAASPAVQAATKVLW
				1		APSPPAAPIPPLRPFGLIQLLCQC
						AQRWQPGSREVKGRLCQARLA
				1		ETSSIAGNLRVQSYERPVDPGV
		ı			1	TMDPLLPGRTIVFAAGDHTLW
		ı				RAPVSSCAQRSGVWSLDNMGV
						NSGSVPFHGVRIQASQLISLSFN
		1				FSFCEADLKTPLHGFLILPAGPK
		1				EEVTVKKEKRERDRDRQREGH
ļ		l				GRGRGRPEVIQSHSIFEQG\PAE
		i				MMKKKGNWDKTVDVSDMGPS
l		1	1		1	HIINIKKEKRETDEETKQILRML
l		1			1	EKDDVDHHALSEHLVTTATFSI
		1			1	GSTGLVVYDYQQVECGGHVRA
		1			1	DLKTQDLYSHAQFGDNNYPGG
					1	VDWYRAGGLQDSDTEDECWS
		1			1	DTEAVPRAPARPREKPLIRSQSL
1		1	1			RVVKRKPPVREGTSRSLKVRTR
		ļ.	22200		504	KKTVPSDVDS
23264	53632	A	23398	297	584	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23265	53633	A	23399	2	149	CMKFGAVTPIGGPPLGDQSPVL
						LLFAL*KRSTYNLGSSDPPAQGI
		1				SHOF
23266	53634	l <sub>A</sub>	23400	1	170	PAGIRHEERSTYDLWSSDRPGQ
						GTSHOF*IIRFOIPWNSGPRLSE*
		1				LLPRSSRLSS
23267	53635	Α	23401	I	1077	
23268	53636	В	23402	289	1359	
23269	53637	Α	23403	3	2233	
23270	53638	Α	23404	1	1491	
23271	53639	В	23405	76	511	
23272	53640	Α	23406	1	2706	
23273	53641	Α	23407	273	862	
23274	53642	Α	23408	1	1080	
23275	53643	Α	23409	134	675	MHYFHSSVQIGLPLTRRSQVLIL
		1				PQPLLPNPIILLSPPLLTPVPAYS
l		1				FIL/PTSPPPPARQFPLKKVAEAK
		1		1		GIVKQLKTDTARSPRKPTGPSQ
				1		TLWVTLTVEDLKHAFFTIPLHP
		1		İ		SSQPLFAFTWTDPDTHQAQQIT
		1		1		RAVLPQVFTDSPHYFSQAQISSS
						SVTYLGIILIKTHVLSLLIMSG
23276	53644	Α	23410	1	895	MEVHKHPFNTLALRHESGCEL
				•		VWAEKHRNQIFPFPKSTASPKQ
	ļ		1			WDTGRHNSDAKQQEPGGREER
		1				NTMEMETKETHFIRGPKTLASV
		1				TDWEGSLPLVFNHCRGTSLIIHP
		1				RFKGVRPRRDACLGPSPLAASP
		1	1			AFLGKGQHALKGLKPVITRLLQ
						HGLLKPINSPFPS\PILPVLKPDK
			i			AYKLVQNLRLINQIVLPIHPVVP
		1				NPYTLLSSIPPSTTHYSVLDLKH
		1				AFFTIPLHPSSQPLFAFT*TDPDT
		1	İ			H*AQQITWAVLPQGFTRQPPLL
						QLSPNFILICYLSRHNSHKNTRA
						LPADRV
23277	53645	Α	23411	1	930	MIQEHADKQVQRLQGVLLGSIP
						TAASRARLHPGEINNHVAHTEP
		1				VWWSLHMDAYEIWCRGSDRR
		ı				TSLGRSIPLPPALCSLRRIHLRPQ
		1				VLRSTSPRNISPISNPGHLSDYTP
1		1				TFQGCQTMQGRLPWSFTLSGK
	1	1	1			SRFSGEGATQRQYPIPQQALKG
1			1			LMPAITRLLQHGLLKPINSPYHS
			l		1	PILPVLKPDKPYKLVQDLRLINQ
		1	l			IVLPIHPMVPNPYTLLSSIPASTT
1		1	1			HYSV\LYVKHAFFTIPLHPSSQP
						LSTFTWTDPDTHQAQQIT*AVL
		1	l	1		PQVFTDSPHYFSQAQISSSSVTY
		1		1		LGIILHENTRALPADHV

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23278	53646	Α	23412	299	1028	RLGSFSSNHPYLGQRQYPIPQQ
		1				ALKGLKPVITCLLQHVLLKPVN
		1		İ		SPYNSPILPVQKLDMPYRLVOD
						LCLINQIVLPIHAMVPNPYTLLT
		1				SIPPSTTHYSVLDLKPAFFTIPLR
		l				LSFQPLFTFSWTDPDTHQAQQI
						TRAVLPQSFTDSPHYFSQAQIS*
		1				RLGSFSSNHPYLGQRQYPIPQQ
		1				ALKGLKPVITCLLQHVLLKPVN
		1				SPYNSPILPVQKLDMPYRLVQD
		l				LCLINQIVLPIHAMVPNPYTLLT
		l				SIPPSTTHYSVLDLKPAFFTIPLR
						LSFQPLFTFSWTDPDTHQAQQI
		l				TRAVLPQSFTDSPHYFSQAQISS
		1				SSVTYLGIPGFAILTKPLYKLTK
		l				GNLADPIDPKSFPHSSFRSLKTA
					1	LETAPILALPVSSQPFSLPTAEV
						WAAQSEFLHKNRDRTE
23279	53647	Α	23413	157	202	RSECARWMEK*CPWL
23280	53648	Α	23414	184	482	EGVSVFLLLSPYQGQPQVSRRC
						S/RQVSQRCPHRGRGPRCPQVS
		ļ				RSHGPSWVGHPGMPLYGS\EQF
						YCGAPGLAGPPNGSRGESLKIS
						VWIFPCRDFGNFN
23281	53649	В	23415	13	862	
23282	53650	A	23416	1	11342	MATRWFYYGHPPRDARVPRIV
	1	1				AEAKEKNLTPGKIIEVDVTIEAQ
		1		ľ		IRQEFLDAERDTGGFVKFFRITK
		1				KKLQEFFLGEEEEQVPHVSPVE
		1				LGAKPTAFNAFIANAVGKTLAV
	1	1				QHASLKAPEPEMTVTHERFYG
		l		l		NYAVWITDVDPRNHRIEVIHVI
						REISGRTSEYVNNLLAQPTPFK
		1		ļ		VFDWLTETQAKEFVRRLNNTQ GVISVYNKLEDGIPVKVAVKAT
						EPTNKEKLEAASTKLKAMGAV
						ATDAYEFHTDLNENE
23283	53651	A	23417	7597	7883	LSSLAWVWQWNHRISVLFHSN
23263	33031	^	23417	1391	/883	SVRTNLVPQFNQLVITESNSCPE
į						FFNOTSAFFKT*VESGCM/CLOD
	ŀ	1				*GIEWFRLRLDVYNSRLVFLTG
		1				ATDYCTG
23284	53652	A	23418	3	1573	
23285	53653	A	23419	6692	6793	GASVAPSIGS*STGNAIPSCLFQ
		Γ.		1		QNVIVPIPQS
23286	53654	Α	23420	1	17727	
23287	53655	Α	23421	6687	6785	VNTGNRTWWEPLFRK*KKTQR
		L				EDTTEKKTKGR
23288	53656	Α	23422	1	2856	
23289	53657	Α	23423	1	7194	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23290	53658	Α	23424	7460	7561	GASVAPSIGS*STGNAIPSCLFQ
						QNVIVPIPQS
23291	53659	A	23425	1468	1857	
23292	53660	A	23426	1	1985	MEPVGIDIIAISWIGAPEYPDNV
		1		1		KRAINAKVEATQKTLQREQEIQ
		1				QRVAEANMEREQAKGVADAIR
		1		1		LRGEALTENPNVMQLEAINKW
		1		i		NGLVPFITSSPGMGKSAIVRQIA
		1				EDFGLKLIDHRLSTSAPEDLSGL
		1				PRFKENGRAEFAPFDELFPLEG
		1				DAVPEGFNGWLLFLDEFNSAR
		1				KEIQAAAYKLVLDRMTGQKKL
		1		l .		HPYVAIVCAGNKATDRAITNNL
1		1				STAMQSRLIHLEMETDFDVFME
		1				DVAIPNKWDERVIAFLNANPNK
1		1		1		LNDFEPDHQEKTFCCSRTWEFV
		İ				NKIVSILPPGPINSEMTLLLAGTI
		1				TSGVATSFVQFTQVYSNMVSL
		1		1	1	NEILDNPKKARMPEDNNLLWA
				l		VVTSLINNTDEDNHSKIFDYVE
		1				RMPFTFKVLYYRSRKTDMSMT
		1		1		DAQLTREYDRCQAKAFMGKTA
		1				AFFGSLLCSLKFRWVKDGCDT
		1				AQTDGEHLEWNPDWFESLLPE
		1				SRVTVLMHELWHVGLLHSVR
		1				MGSRDPEVWNYACDIYINNQLI
İ		1				QDGYSFIGIENCWKDPKYAGW
l				ŀ		AEEQIYHDLMSKHPRPPKASGA
		1				FGTGSGGDMKPSTSKASLKLSG
		1		1		GSLPGVLPGKMEEVITQFLKPV
i		1				VPWQVLLERFFNDLQETYYSW
		1				QRPNRRYPDMYLPSSMDDDCR
		1				LEHLAYFL*NEADLTGLLVLK*
						TDDGQSVSGQNLLQSDEQLQIY
23293	53661	Α	23427	2365	2784	
23294	53662	Α	23428	5789	5980	DQNMRFDNTFETYYWKPTIIVII
		1		i	i	IRKLLMNDSSKDTLK*LRCWK
						NWLQILKKHTIHFIMPLQL
23295	53663	Α	23429	I	6819	
23296	53664	Α	23430	376	2052	SKVCGNDGFADDLKHLLIFTGA
		1				QSGEDIVPFQLEDGHSLGEVVL
		1	l		1	LHGGSGVESGQR/CGRTSAGSC
		ı				Y*SPGKICCCNLLVTKTEGFFAP
						VPIQLENTNQVTNCSDNW
23297	53665	Α	23431	2	2189	
23298	53666	Α	23432	2	1583	
23299	53667	Α	23433	2142	2242	GASVAPSIGS*STGNAIPSCLFQ
		1_				QNVIVPIPQS
23300	53668	Α	23434	3	1188	
23301	53669	Α	23435	470	862	
23302	53670	A	23436	1	1352	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
23303	53671	A	23437	1837	1955	SKCSIGLYQRYPYSNGC*NSGCI CRVY*CCREHRGYVRN
23304	53672	A	23438	1447	1565	SKCSIGLYQMYPYSNGC*NSGC ICRVY*CCREHRGYVRN
23305	53673	Α	23439	773	823	
23306	53674	Α	23440	3	1529	
23307	53675	Α	23441	1	1794	
23308	53676	A	23442	236	427	DQNMRFDNTFETYYWKPTIIVII IRKLLMNDSSKDTLK*LRCWK NWLQILKKHTIHFIMPLQL
23309	53677	A	23443	8510	8611	GASVAPSIGS*STGNAIPSCLFQ QNVIVPIPQS
23310	53678	A	23444	516	669	ATSELKRKASLGLMEPDVSRK* ARCSSLPSSSIGDGRYISGYLRL GRCHE
23311	53679	A	23445	410	593	FLNLIQPQGKRKLLLKRCRSLLP KN*CQHSKQHKLSISSVVILNIT NS*PNKVKLMPILAM
23312	53680	В	23446	1	2286	
23313	53681	Α	23447	1418	1723	CNTNIIFNCHFPTKLGCICKAMK VVSPLIHNSIVILLNQITIDFFSTL AQICTFICLTYSKQTTTRIASFK KVIEEFLSLAPIEKVELKEHS*YI SISEKV
23314	53682	В	23448	1	6167	
23315	53683	Α	23449	1	3199	
23316	53684	Α	23450	1	5733	
23317	53685	Α	23451	709	1099	
23318	53686	A	23452	5039	5230	DQNMRFDNTFETYYWKPTIIVII IRKLLMNDSSKDTLK*LRCWK NWLQILKKHTIHFIMPLQL
23319	53687	Α	23453	664	993	
23320	53688	A	23454		19859	MASIDSIAVCNPRQVYKFVEHC IRSGLVPFITSSPGMGKSAIVRQI AEDFGLKLIDHRLSTSAPEDLSG LPRFKENGRAEFAPFDELFPLEG DAVPEGFNGWLLFLDEFNSAR KEIQAAAYKLVLDRMTGQKKL HPVVAIVCAGNKATDRAITNNI STAMQSRLIHLEMETDFDVFME DVAIPNKWDERVIAFLNANPNK LNDFEPDHQEKTFCCSRTWEFV NKIVSILPPGPINSEMTLLLAGTI TSGVATSFV
23321	53689	В	23455	74	240	
23322	53690	Α	23456	143	384	
23323	53691	Α	23457	1	3267	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide scquence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \-possible nucleotide insertion)
23324	53692	A	23458	2	550	KRSSEHIQRIDRDVSGTLRKHIF FRDRYGTKQRELLHILLAYEEY NPEVGYCRDLSHIAALFLLYLP EEDAFWALVQLLASERHSLQG FHSPNGGTVQGLQDQGEHVVA TSQPKTMGHQDKKDLCGQCSP LGCLIRILIDGISLGLTLR'LWDV YLVEGEQAFMPITRIAFKVQQK RLTKYSR
23325	53693	Ā	23459	72	630	
23326	53694	A	23460	I	2835	MDALMLALGALMILAGLOD KSAQVSDPDGAEPGPGRATGT DPLVQIIIWKLGISEALSIPGGRL YRDLWSLTPVCLHIPGIAHHGP FTLGRMDVVEVAGSWWAQER EDIIMKVEKGHRAGLPEDKGPK SFGSYNNNVDHLGIVHETELPP LTAREVKQIRREISRKSKWVKM LGEWDTYKNSKKLIDRAYQGIP MNIRGPMWSVLLNIEEIKLKNP GRYQVRSARAQPTGQAVSGAQ VSSWRERQDHPGELGVKI

3075

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	tocation of first	codon for fast amino acid	*=Stop eodon, /=possible nucleotide
	sequence	ı	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
23327	53695	A	23461	li .	832	MGWRARWLMPVISALWEAED
		1		1		PIPPELTRLLQQSQDPLLMGLFP
		l				TNPKEKTQEEPPGQSRAPVLTV
		1				VSKFKVLAGLLHLGNIQFAASE
		1				DEAQPCQPMDDAKCEGQGVG
ĺ		l				WAGPVRKASLEQLLQVLHSTT
		l				PHYIRCIKPNSOGOAOTFLOEE
						AGAGRRRORRROKOORRRRRO
		l				LGGGGDGGNGSVGDDGATRGS
		1				RGCALEKECVATPRRGLLFSRP
		l				HKLAFAVGGSVTSVGPKPRAP
		l				MDVVEVAGSWWAQEREDIIM
		l				KYEKGHRAGLPEDKGPKPFRS
1		l				YNNNVDHLGIVQDFADFGTTIK
1		1				QDFRLLGQTSVDRLLQLSQGQ
ļ		i				AVKGNQLLPVSLVKRKTTLAP
1				1		NTQTASPRALADSLMQLARQV
		l		1		SRLESGHFYDAHNPFLPPPSMS
		1				KQAKKTGVPSEGLCRPGGEGQ
		1		1		SAVARLTGEKKNHPGAQYANR
		l				LSPRVGRFINAAGTTGFPTGKR
1	l	1				AVSATQLIGLQLILLGEVLQKV
		1				LTASYPAQSSVPLTSGPVRQIM
1		1		i		KEKGKRSSEHIQRIDRDISGTLR
i		1				KHMFFRDRYGTKQRELLHILLA
		1		ļ		YEEYNPEVGYCMDLSHIAALFL
		1				LYLPEEDAFWALVQLLASERHS
	ļ.	l				LQAARAPAAIGAHERADQAQIS
						LGLTLRLWDVYLVEGEQALMP
		l				ITRIAFKVQQKRLTKTSRCGPW
1		l				ARFCNRFVDTWARDEDTVLKH
		-				LRASMKKLTRKQGDLPPPAKPE
23328	53696	A	23462	3	497	SSECLTMAWIPLLIPLLILCTVS
						VASYELTQPASVSVSPGQTASIT
1						CSGDVLAKKKYARWFQQKPG
						QAPVLVIYKDSERPSGIPERFSG
		1				SSSGTTVTLTISGAQVEDEADY
		1				YCYSAADNNGWVFGGG\TKLT\
		1				VLSQPKAAP\SVTLFP\PSSEELQ
23329	52607	١.	22462	1	393	ANKAHTGVSQ
23329	53697	A	23463	]'	393	FRSGSELVKAGLRAFFENAAED
		1				LEKTSENLKLGKFTHSRTQIKG
		1				VSQNINYTTVALLPILTSIFEHV
		1				TQHQFGMDLLLGDVQISCYHIL CSLYSLGTGKNIYVE\ILGMPDT
		1				
23330	53698	Α	23464	319	508	VEDMCPDIPQLECLMKEIN
23331	53699	C	23465	94	222	,
23332	53700	A	23466	1	636	
23333	53701	Â	23467	1	483	
233334	53702	Â	23468	1	1797	
~5554	100/02	10	22700	ı·	1.121	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23335	53703	Α	23469	l	4275	
23336	53704	Α	23470	l	1848	
23337	53705	Α	23471	122	14928	ARSWRRRWRTRTKRRRKSAA
		1				AVSARRAAAGGSRGAGGWGT
						ADASGAMAEGGEGGEDEIQFL
1						RTEDEVVLQCIATIHKEQRKFC
						LAAEGLGNRLCFLEPTSEAKYIF
						PDLCVCNFVLEQSLSVRALQEM
						LANTGENGGEGAAQGGGHRTL
		1				LYGHAVLLRHSFSGMYLTCLTT
						SRSQTDKLAFDVGLREHATGE ACWWTIHPASKQRSEGEKVRIG
		l		l		DDLILVSVSSERYLHLSVSNGNI
	İ	1				OVDASFMOTLWNVHPTC
23338	53706	A	23472	134	1063	RPPGLTRSPSSPAPARSPTRRQS
23330	33 /00	^	23472	134	1003	TRRRSGATTSCSQRLWATWNG
						CD/VCLNQSLREIPTDDKGFTAI
l		l				HFAAQWGKLACLQVLVEEYKF
						PVDLLTNNSQTPLHLVIHRDNT
						TVALPCIYYLLEKGADLNAOTO
	1					NGSTPLHLAARDGLLDCVKVL
		l				VOSGANVHAQDAMGYKPIDFC
		l				KIWNHRACARFLKDAMWKKD
						KKDFAREMTKMKMFKSQLTL
		1				MEHNYLIEYQKEHKILREAAIR
		1				KWLHGKLHPGHSLVSNTKQAR
						ATALSKTPEQRESQRSRSFHPSV
	1					DARLQCIPQPTEMPKPIYRKPTV
						KRPHNVEC
23339	53707	Α	23473	1	1280	MAGLGFWGHPAGPLLLLLLLV
					1	LPPRALPEGPLVFVALVFRHGD
1					1	RAPLASYPMDPHKEVASTLWP
ļ						RGLGQLTTEGVRQQLELGRFLR
		1				SRYEAFLSPEYRREEVYIRSTDF
		1				DRTLESAQANLAGLFPEAAPGS PEARWRPIPVHTVPVAEDKLLR
		l				FPMRSCPRYHELLREATEAAEY
		l			j	QEALEGWTGFLSRLENFTGLSL
					1	VGEPLRRAWKVLDTLMCQQA
	1					HGLPLPA WASPD VLRTLAQISA
					1	LDIGAHVGPPRAAEKAQLTGGI
	1	1				LLNAILANFSRVORLGLPLKMV
1	1	1			1	MYSAHDSTLLALQGALGLYDG
	1				1	HTPPYAACLGFEFRKHLGNPAK
		1			1	DGGNVTVSLFYRNDSAHLPLPL
		1			1	SLPGCPAP\CPLGRFYQLTAPAR
	1	1			1	PPAHGVTCHGPYEAAIPPAPVV
	1	1			1	PLLAGAVAVLVALSLGLGLLA
1	1	1			1	WRPGCLRALGGPV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23340	53708	A	23474	133	672	AVGWESGK WDSGRGQGGGQE
						OMWSCOSDMLYPPDOVGAWG
	l					LPGPTGPKGDAGSRGPMGMRG
	i i	l				PPGECPQCCPTAPLAIPSTRWAL
l		l				PCSWPPGPPGPPGPPAPVGPPHA
		1				\GSPSMVSPPGIPAGGGGGGVAI
	Į.					STVPATIKTRROEVSPELPRP/PG
		1				PAPPPGPDVRPWLLRPPGPMGK
						LSPGQG
23341	53709	Α	23475	135	677	AVGWESGKWDSGWGQGGGQE
	i	1				QMWSCQSDMLYPPDQVGAWG
						LPGPTGPKGDAGSRGPMGMRG
		1				PPGECPQCCPTAPLAIPSTRWAL
						PCSGPPGPPGPPGPPAPVGPPHA
		l		ľ		\GSPSMVSPPGIPAGGGGGGVAI
ŀ		1				STVPATKKTRRQEVSPELPRP/P
ļ						GPAPPPGPDVRPWLLGPPGPMG
				1		KLSPGQG
23342	53710	Α	23476	220	745	TVIQRTGLHSVGPGDPQQGLRE
l		1				AKLGTMVRLVLLPTPRRTTEVR
						TGPERELSMVLEAILGSLLETAC
	İ	1		1		DHRQTPPSSKPQA/GSSRLREGT
						SAGQPCS/LRLRLGPRAGGRRC
	Į.	1				CRDIERSSSSSVCGRKCKRKKR
		l				KRGMCQSRHKHGCRRPQEGPR
						CNDGRALCSPRPQTTPGPLSTA
23343	53711	A	23477	1	2433	
23344	53712	A	23478	3	803	FPGRRFRVRACVAGGSHCLFNS
	1			ļ		CCCCTSASLPARERHEWPGPAA
	1	1				AAAAAAAARRSQTQREGLFA
		1			l	GWGGGFAMSDDDSRASTSSSSS
						SSSNQQTEKETNTPKKKESKVS
		1				MSKNSKLL\SPRAKGIQKEL\AD
		1				ITLDPPPNCSAGPKGDNIYEWR
		1				STILGPPGSVYEGGVFFLDITFTP
						EYPFKPPKVTFRTRIYHCNINSQ
			1			\GVICLDILKDNWSPALTISKVL
			1		1	LSICSLLTDCNPADPLVGSIATQ
				1		YMTNRAEHDRMARQWTKRYA
	1	1				T

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino neid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23345	53713	A	23479	372	1839	GEAGGTEAEPENGAGGPRVPG WWGWSSTSSPRGCWDSGRSAA AIMDELWDKRTGLAARKMPHP RRYHSSERGSRGSYCEHYRS/R KYHKQRRSRSWSSSSDRTRRRR REDSYHYRRCSRTSRSSSQH SSRKAKSVEDDTEGHILIYHVGD WLGERYEIVSTLGKGTFGRV QCVDHRRRGARYALKIIKNVE KYKEAARLEIKYLEKINEKDPG KNLCYQMFDWFDYHCHMCISL ELLGLSTEDFLKDNNHLPYPH UYHHMASQLCQAVKFLHDNK LTHTDLKPENILFVNSDVELTY NLEKKRHERSVKSTAVRYODF SAATFDHEHISTIVSTRHYRAP EVILELGWSQPCDVWSIGCIIFE RILGWSQCDVWSIGCIIFE RILGWSQCDVWSQCDVWSIGCIIFE RILGWSQCDVWSQCDVWSIGCIIFE RILGWSQCDVWSQCDVWSIGCIIFE RILGWSQCDVWSQC
23346	53714	A	23480	28	941	QSARPSLKRARSQRGRPJPSRA LVFLEFFFFFFNFLPHITQPSAH LVFLEFFFFFNFLPHITQPSAH KOMITNAGPLHPYGPGHLNRLD NFVPNDRPTWAVYGGRAGCEPI GDFGRATVPCCWVLQCVGFH LVIEGLVPFSNYERFCFGOQSLL NKLWKEYGQGETARLHPGG DKFHSVALGTITSFALWGEPFSL WVVIAFLLQHPLNFFLIQLVVS VGQNYGGDVFVFLKEHFRDS KHGKPGPTPLLLPGFSFVFHEM PLWLVPAPGSPLCLDAVKAPS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
23347	53715	A	23481	1	1386	MWRNLKLPRDLLNGFDONADS
23347	33713	ľ`	23107		1.500	DMDNEVOAEVVTDGDEELIGN
					·	WSKGHSCYILAKRLAAFFPCPK
						DLWKFELERDDLGYLAEEISSS
						EAFKSVSYEKAHAWNPLESTC
						RHASLALAVVLORRDWENP\G
						VTOL\NRLAAHPPFASW\RNSEE
						ARTDRPSQQLRSLNGEWRLMR
						YFLLTHLCETLVKVKDAEDOL
			1			GARVCVLEROIFDFLGYOWAPI
1			ĺ			LANFVHILIVILGLEGTIOYRPRY
						ITGILSFPKHQESLSLKFTCLTEA
						ERDTDALLHLENACRESTATOR
						IQGFFLKDKKRKKQQPQQQKQ
						OOWAKVOLGPWLORVOAPSL
						GSFHGVLSLRVHRSKELRFRNC
			ŀ			CLDFRGCMEMPGCPGRGVLQA
		l				ONPRGEPLLGOCRREMCGGSP
						YIVPTGALPSGAVRRRPPSSRPH
			i			NGRPTDSLHCVPGKAADTQHQ
						PLKAARRAAVPCKATGSELPK
23348	53716	Α	23482	3	732	SPLTPTATSGNPSKASSPGGLGS
						LVSTASSLPAGEKNSPRTGPGP
						RRAG\AGIPNSGPDTVEGPYNL
			ł			CGRQGRWGGLIISLASEMKLQT
						LQTSVTTAKATLLELFVPLGGL
						VASLASEVRLQIFPRGGHLQSA
			ŀ			KCGLLLHFSLPTSRSGRRKANR
						PMAVTVLLVLIRDCADLLAVK
						AVIHCYQDQWPLRGSSRPANTP
						PPKASVSRRPPGHRGVRRGSPL
						LQVGARRVGAEAPALLELQSPS
	53717	A	23483	1	234	
23350	53718	Α	23484	1	256	MLNLQGKEWLELEEDAQKAYI
						MGLLDRLEVVSRERRLKVARA
		1				VLYLAQDGEKFCPKRDCTNNS/
						RY/NSI*MIQMMRFGTLELMRL
	53719	Α	23485	1	3333	
	53720	Α	23486	1	2661	
23353	53721	Α	23488	3	433	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon fur last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22254	62720	<u> </u>	02400	3	990	EKMNTADQARVGPADDGPAPS
23354	53722	A	23489	,	1990	,
1		l			ł	GEEGEGGGEAGGKEPAADAA
1		l				PGPSAAFRLMVTRREPAVKLQ
1		l				YAVSGLEPLAWSEDHRVSVST
1		1				ARSIAVLELICDVHNPGQDLVIH
		l				RTSVPAPLNSCLLKVGSKTEVA
		l				ECKEKFAASKDPTVSQTFMLDR
		l				VFNPEGKALPPMRGFKYTSWSP
		l				MGCDANGRCLLAALTMDNRL
		l				TIQANLNRLQWVQLVDLTEIYG
		1	i			ERLYETSYRLSKNEAPEGNLGD
		1				FAEFQRRHSMQTPVRMEWSGI
		1				CTTQQVKHNNESRDVGSVLLA
		1				VLFENGNIAVWQFQLPFVGKES
					1	ISSCNTIESGITSPSVLFWWEYE
					į.	HNNRKMSGLIVGSAFGPIKILPV
					l	NLKAVKGYFTLRQPVILWKEM
		l				DQLPVHSIKCVPLYHPYQKCSC
						SLVVAARGSYVFWRLLLISKAG
		1			1	LNLHNSHVTGLHSLPIVSMTAD
		l				KQNGTVYTCSSDGKVRQVIPIF
		l				TDVALKFEHQLIKLSDVFGSVR
1		l				THGIAVKPCGAYLAIITTEGMIN
1		l			İ	GLHPVNKNYQVQFVTLKTFEE
		l				AAAQLLESSVQNLFKQVDLIDL
1		l				VRWKILKDKHIPQFLQEALEKK
						IESSGVTYFWRFKLFLLRILYQS
						MOKTPSEALWKPTHEDSKILLV
		1	ľ			
						DSPG\MGNADDEQQEEGTSSKQ
		1				VVKQGLQERSKEGDVEEPTD\D
		l				SLPTTG\DAG\GREPMGRRNSW
		<u> </u>				GNSKGKIRSCWEMRLTREHMK
23355	53723	A	23490	1218	1446	RCTDSQV\RSPTPLHLAAQACSL
		l		ŀ		ETTVCLLCSKADYTLSEKRGW
		l				MPIHFAAFYDNVCIIIALCRKDP
				L		SLLEAEATAE
23356	53724	A	23491	2	632	NMAKTYDYLFKLLLIGDSGVG
		l				KTCVL\FRFSE\DAFNSTFIFTIRI
		l				DFKIRTIELDGKRIKLQIWDTAG
	İ	1				QERFRTITTAYYRGAMGIMLVY
		1				DITNEKSFDNIRNWIRNTEEHAS
		l				ADVEKMILGNKCDVNDKRQVS
	1	1	l			KGS/GENKLALDYGIKFMETSA
	1	ı	l	l		KANINVENAFFTLARDIKAKMD
1		ı	l	l	l	KKLDGNSP\QGSNQGVKITPDQ
		l		-		OKRSSFFRCVLL
$\overline{}$	L	Ц.				Z.C.CO. I IC VED

NO:   of peptide   sequence   s	SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
23357   53725   A   23492   I   I018   MVFYHVGQAGLELLTSGPHCP DSQAADKAESATLLSRARLGA YCACASLERRWHFPSGRECN MAKTYDVLKKLLUBIOSGVGK TCVLFRFSEDAFNSTFISTIGIDF KIRTILELDGKRIKLQIWDTAGQ ERRTDHNRPTYRGAMGIMIL VYDITHEUSSFDONIRNIWIRNIE EHASADWEKMILGONLCOWND KRQVSKERGEKLALDIYGIKFM ETSAKANINVENAFTLLARDIK AKMDKKLGPARPHGTLLGKGE EFAGVAGASASGASLGTLFKQ TRPCPVNPQASREHTASTIPLKG QKDMMEGNSPQGSNQGVKITP DQKRSSFFRCVLL   1392   130	1	sequence	l	09/540,217		of peptide sequence	deletion, \-possible nucleotide insertion)
DSQAADKAESATLLSRARLGA			l		sequence		
DSQAADKAÈSATLLSRARIGA	23357	53725	A	23492	1	1018	MVFYHVGOAGLELLTSGPHCP
VCACASLERRWPHFPGRECN   MAKTYDYLFKLLLIGDSGVGK   TCVLFRSEDAPNSTFISTIGIDF   KIRTIELDGKRIKU,QIWDTAGQ   ERFRTDHNRPTYRAMGAMGML   VYDITHEKSFIDNIRNWRINE   EHASADIVEKIMILGNSTOWND   KRQVSKERGEKLALDYGIKFM   ETSAKANINYENAFFILARDIK   AKMDKKLGPARPHGTLLGKGE   EFAGVAGASASGASLGTLFKQ   TRPCPVIPQASRETHTASTIPLK   GKDNIMGNSPQGSNQGVKITP   DQQKRSSFFRCVLL			1		ľ		
MAKTYDYLFKLLUIGDSGVGK	ł		l		İ		
TCVLFRFSEDAFNSTFISTIGIDF   KIRTILELDGKRIKLQIWDTAGQ   ERRRTDHNRPTYRGAMGIML   VYDITHEUKSFIDNIRNWIRNIE   EHASADWEKMILGNICDVND   KRQVSKERGEKLALDIYGIKFM   ETSAKANINVENAFTILARDIK   AKMDKKLGPARPHGTILLGKGE   EFAGVAGASASGASLGTLFKQ   TRPCPVNIPQASREHTASTIPLKG   GKDMMEGNSPQGSNQGVKITP   DQKRSSFFRCVLL			l	1			
RIRTIELDGKRIKLQIWDTAGQ			l		l		
			i				
VYDITHEIKSFIDNIRNIWIRNIE			i				
EHASADIVEKIMILGNIKCDVND   KRQVSKERGEKLALDYGIKFM   ETSAKANINVENAFTLARDIK   AKMDKKLJPARPHGTILGKGE   EFAGVAGASASGASAGATIPKQ   TRPCPVNPQASREHTAFSIPLKG   QKDMMEGINSPQGSNQGVKITP   DQKRSSFFRCVLL			l				
RRQVSKERGEKLALDIYGIKFM   ETSAKANINYENAFTILADIK   AKMOKKLGPARPHGTILLGKGE   EFAGVAGASASGASLGTLPRQ   TRPCPVNPQASRETHATSIPLKAG   QKDNMEGNSPQGSNQGVKITP   QQKRSSFFRCVLL			l				
ETSAKANINVENAFFILARDIK   AKMKKLGPARPHGTILGGE   EFAGVAGASASGSASLGTLPKQ   TRPCPVNPQASREHTAFSIPLKG   QKDNMEGNSPQGSNQGVKITD   DQQKRSSFFRCVLL			l			l	
AKMDKKLGPARPHGTILGKGE			l				
EFAGVAGASASGASLGTLPKQ TRPCPVNPQASREHTAFSIPLKG QKDMMEGNSPGQSSNGGVLITP DQQKRSSFFRCVLL			l			İ	
TRPCPVNPQASREHTAFSIPLKG   QKDNMEGNSPQGSNQGVKITP   QKDNMEGNSPQGSNQGVKITP   QQKRSSFFRCVLL			i				
QKDNMEGNSPQGSNQGVKITP			l				
DQQKRSSFFRCVLL			l				
23358   53726			l				
23369   53727   A   23494   I   1392	23358	53726	Ι	23493	1	130	DQQRRSSITRE VEE
23360   53728   A   23495   445   649   YPSLTNSKLLVHHFMELCWD KCVEKPQGIALDSRENCLSSC WDRFIDTILATISFR-QUVOKG QQ   VIVAPGKKAQRTQK/APKTKTK KPHLSPCPSP/WRSRSCQPMQQS WSGAQVHIPQPPHTTCITWA   VIVAPGKKAQRTQK/APKTKTK KPHLSPCPSP/WRSRSCQPMQQS WSGAQVHIPQPPHTTCITWA   QASSSPISSVTCKALPPGQTK KGLVASQPEPVSMGTSSPTPS LAPEHSSRASLEPSSSA*PSLAT* SRAGGSQA   QASSSPISSVTCKALPPGQTK GGLVASQPEPVSMGTSSPTPS LAPEHSSRASLEPSSSA*PSLAT* SRAGGSQA   QAMHGRSPGVTVRASGLRNEG DGPLGSVPTQARATGAPGHSAT PSPAWRAW/SSSSKAS/WLPDA AGPPTPPWSGFSSPTSW   QAMFGSSPTSSW   QAMFGSSPTSSWASWSSSKAS/WLPDA AGPPTPPWSGFSSPTSW   QAMFGSSPTSSWASWSSSKAS/WLPDA AGPPTPPWSGFSSPTSW   QAMFGST   QAMFGST   QAMFGST   QAMF			-		-		
XCVEKPQGIALDSRTENCLSISC   VIDRIBITYLAITSRFAQIVQKG   QQ   QQ   QQ   QQ   QQ   QQ   QQ					<u> </u>		YPSLTNSKLLVHH\FMELC\WD
VORFIDITALATISRFAQIVQKG   QQ   Q23361   53729   A   23496   210   397   VIYAPGKKAQRTQK/APKTKTK   KPHLSPC'SPIWRSRSCQPMQQS   WSGAQVHIPQPPHTCITWA   QASSSPISSVICKKALPPGQTK   KGLVASGDPEPVSMGTISSPITS   LAPPHSSRASLEPSSA*PSLAT*   SRAGGSQA   S3731   A   23498   780   1022   QAMHGRSPGVTVRASGLRNEG   DGPLGSVFTQARATGAPGHSAT   PSPAWRAWSSSSKASWLPDA   AGPPTPPWSGPSPTSW   QAMPPTPWSGPSPTSW   QAMPPWSGPSPTSW   QAMPPTPWSGPST   QAMPPTPWSGPST   QAMPPTPWSGPST   QAMPPTPWSGPST   QAMPPTPWSGPST   QAMPPTPWSGPST   QAMPPTPWSGPST   QAMPPTPWSGPST   QAMPPTPWSGPST   QAMPPTPWSGPST   QAMPPTPWS			ľ			1	
CQ   CYPAPGKKAQRTQK/APKTKTK   KPHLSPCPSP/WRSRSCQPMQQS   WSGAQVHIPQPHITCITWA			l		l	ľ	
23361   53729			l				
KPHLSPCPSP/WRSRSCQPMQOS   KPHLSPCPSP/WRSRSCQPMQOS   WSGAQVHIPQPHITTCITWA	23361	53729	A	23496	210	397	
WSGAQVHIPQPHTTCITWA			Γ.				
23362   233730			ı				
KGLVASGDFEPVSMGTSSPTPS	23362	53730	A	23497	55	286	
LAPEHSSRASLEPSSSA*PSLAT*   SRAGGSQA			l				KGLVASGDPEPVSMGTSSPTPS
SRAGGSQA			l				
23363   53731   A   23498   780   1022   QAMHGRSPGVTVRASGLRNEG   DGPLGSVPTQARATGAPGHSAT   PSPAWRAW\SSSSKAS\WLPDA   AGPPTPPWSGPSPPTSW   23364   53732   A   23499   I   782			ı				
DGPLGSVPTQARATGAPGHSAT   PSPAWRAWISSSSKASWLPDA   AGPPTPPWSGPSPPTSW   23364   53732   A   23499   I   782	23363	53731	A	23498	780	1022	
PSPAWRAWISSSSKASWLPDA   AGPPTPPWSGPSPPTSW   1   782   1   782   1   1   1   1   1   1   1   1   1							
23364 53732 A 23499 I 782			I	l	l		
23364 53732 A 23499 I 782	1		ı		i		AGPPTPPWSGPSPPTSW
23365 53733 A 23500 275 688	23364	53732	Ā	23499	I	782	
	23365	53733	A	23500	275	688	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ı	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
1		l		sequence		
		<u></u>				
23366	53734	A	23501	1166	2271	LRCILTLISRWCS*VSTVANLPP
		l				GISKWGANWQRCEMKE*CWSP
1		1			i	AVTWCITCAQRSG/IGDSSPYPG
		l				ATSFNEYVKANLTWRGPVEQH
		1				PLVNYLDHERGTLSNPTARHYL
1		l				PVLYVLRAILWARTPLPIPGRTT
1	1	l	:			VSKVKMKKRILRAMRQKQQVL
i		l				PLLVGLGLDELSMSAPSIPAAK
		l				ARMAOLDSRECRKLLNOAMA
	l	l				CRTSLEVEHLLAQFRMTQQDA
		1		i		PLVTAECITLESDWRSKEEVLK
		1				GMTDNLLLAGRCRYPRKLEAD
	ļ	l				LWAREAVFSTGLGFSFAIPHSK
	ļ	l			1	SEHIEQSTISVARLQAPVRWGD
		İ			1	DEAOFIIMLTLNKHAAGDOHM
		l				RIFSRLARRIMHEEFRNALVNA
		l				ASADAIASLLNMNWTVKEHHE
23367	53735	A	23502	1	2604	ASADAIASEENWINWTVKEITIE
23368	53736	A	23502	306	468	TRHCDRROPRRGHRRAGT/CNO
23300	33730	l^	25505	300	1700	I*KGCCCVRTKQPLSSVLISPNF
i		1				CPCSSFLKNS
23369	53737	A	23504	102	509	AAQQGTMARSLLLPLQILLLSL
23303	33131	<u>۱</u> ^	23304	102	1309	ALETAGEEAQGDKIIDGAPCAR
1	1	1				GSHPWQVALLSGNOLHCGGVL
1		1				VNERWYLTAAHCKMNEYTYH
		1				LGSDTL/E/ADRKAQRIKPSKSF
		l				RHPGYFTOTHVNDLMLVKLNS
		l				OARLSSMVK
23370	53738	A	23505		1032	MMQATVGFEDGDAATSQVSPS
23370	23/36	Ι^	23303	1	1032	VNKVHSSIYFLDFFEEYIVIAFKI
		l				
		ı			Į.	LRRGPDISCSQIINGEDCSPHSQP
I		1			l	WQAALVMENE/LFCSGVLVHP
ł		l l			ľ	QWVLSAAPVSRKYNRPLLAND
1		1				LMLIKLDESVSESDTIRSISIASQ
1		l			ł	CPTAGNSCLVSGWGLLANDPT
		l				SR\LPPPLECRNPVPQLLLPPEKP
ł		1				RTAPDTLGSSPVPKSRTGSSGSS
1					İ	FLLPAPPSPFPRPHHGTPPTSCG
1						QDVDVPALAGGSLGSPAGEAL
1		1			i	PRLAAPERLSRHHGKIPSPAPAD
		1	l			LTAILALETAGEEVPFPLTSCA
l		l	1		l	WMSSSSPPGLHEGLQGLTGKFH
1						AVRWHPRLQEKRLQSVPSPDSR
		1				IPGARRPAAR
23371	53739	Α	23506	303	393	
$\overline{}$		_			<del></del>	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \≈possible nucleotide inscrtion)
				sequence		
23372	53740	A	23507	391	1170	GSFFFFFKYFAYSPMPKNSKVV
23312	33740	Α.	23307	391	1170	KRELDDDVTESVKDLLSNEDA
				i		ADDAFKTSELIVDGQEEKDTDV
						EGK\SEVEDERPAWNSKLQYIL
						AQVGFSVGLGNVWRFPYLCQK
		l				NGGVVEPECEQSSATTYYWYR
			ľ			EALNISSSISESGGLNWKMTICL
		ŀ				LAAWVMVCLAMIKGIQSSGKV
		ŀ				NTYDRVTMSATVLALQEQWQT
						FSPIRSGNGHLTGSGAQRTPCRI
						RRGGSQLRVCDGGKQQWWTV
		_				SQSSARAITNTDQKSVQLQDLIE
23373	53741	Α	23508	45	546	EGGGSRSLWFRELIGQPVQSFIE
						PVAAGGAVALNVQ/RSCGVVS
						AAPACL*SRPHSWRWGGPACW
						RTPGG/LASRSSSSASILMSSSRA
			ļ			SFTRSLSLLSTHKDQGLVCSGSS
		1	l	ĺ		GATGAGSC/LWPPTSHTVKLMF
		1				LYSTVSTLKPMVGMVVTISPSL
1		l				SL*RIVVFPAASRPT/YEDAHFFF
						CQKGP*RGSWSSLQHPGPP
23374	53742	A	23509	1367	2142	VRRPQGCLGKRSLLSRVCALVP
		1				WGIWP/WSLAVGPQYSSLGSQP
		l				ILCASIPGLVPKQLRFCRNYVEI
						MPSVAEGIKIGIQECQHQFRGR
						RWNCTTVHDSLAIFGPVLDKG
						YFGYLETITVPSDSVVSTKAETP
						DKGVCGRELGFRGGWLPELQS
						LDEISMGWALCSRLQIKATRES
		l				AFVHAIASAGVAFAVTRSCAEG
						TAAICGCSSRHQGSPGKGWKW
		1				GGCSEDIEFGGMVSREFADARE
						NRPDARSAMNRHNNEAGROV
					1	GSPPARVLGKKEPPOOGVCPGS
						LGHMARSLAVGPQYSSLGSQPI
						LCASIPGLVPKQLRFCRNYVEI
			ŀ			MPSVAEGIKIGIQECQHQFRGR
			l	l		RWNCTTVHDSLAIFGPVLDKG
				l		YFGYLETITVPSDSVVSTKAETP
			I	l		DKGVCGRELGFRGGWLPELQS
1			l	l		LDEISMGWALCSRLOIKATRES
						AFVHAIASAGVAFAVTRSCAEG
		1	l	l		TAAICGCSSRHOGSPGKGWKW
		1		l		
			1	I		GGCSEDIEFGGMVSREFADARE
			l	L		NRPDARSAMNRHNNEAGRQ

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23375	53743	Α	23510	2094	2619	ASPASPAPCSGIPRT*SPYPSPSR
		l				TQ\APRARGPAIHAHPGP/RGCT
			ŀ			ALPROGPPTPGSARAPPRIPAHR
						APTSPTLNFCLASSRQSMPPAPR
						PPPVPATNPVTPGCGRPPPTLGC
						RNPQTLRPPPGLPSPALLRRPLP
			ł			PPARVGMPPHQPPLPTTPACWR
						PVDWARAEGAGRAVC
23376	53744	Α	23511	3050	3317	PARTSPSASGPP**DLISALWTPS
						QPRAPTRQSLNSLSSSAPSSSTIG
						PSPSPPAPGPAPPLPWLPPSPSSL
						PRLSQVSRRPHSHLHPQ
23377	53745	A	23512	862	2954	AEHTLLVCFLNTSLTTTSTHVES
		l				FGGQLRGSQIGHPRRPNKMPYA
			l			WSIPQDPPLKPERDPGGPLGVS
		l				LSLPVNQYGESEEPMGDDDYD
					Į.	EYSKELNQYRRSKDSRGRGLSR
						GRGRGSRGRGKGMGRGRGRG
1						GSRGGMNKGGMNDDEDFYDE
1			İ	}	l	DMGDGGGGSYRSRDHDKPHQ
İ						QSDKKGKVICKYFVEGRCTWL
l						YHTTGNCINGDDCMFSHDPLTE
						ETRELLDKVEMTGSSVFDYIHP
			İ			GDHSEVLEQLGLRTPTPGPPTPP
					i	SVSSSSSSSSSLADTPEIEASLTK
						VPPSSLVQERSFFVRMKSTLTK
						RGLHVKASGYKVIHVTGRLRA
						HALGLVALGHTLPPAPLAELPL
						HGHMIVFRLSLGLTILACESRVS
						DHMDLGPSELVGRSCYQFVHG
						QDATRIRQSHVDCETHLHPPSL
						PTTPQTRASHSLVPGSPSSGLPT
						VPPAAPTAPGTPFSIGRAARRLS
						DASGLLDKGQVMTGYYRWLQ
			İ			RAGGFVWLQSVATVAGSGKSP
			1			GEHHVLWVSHVLSQAEGGQTP
		1	ĺ		I	LDAFQLPASVACEEASSPGPEPT
						EPEPPTEGKQAAPAENEAPQTQ
						GKRIKVEPGPRETKGSEDSGDE
						DPSSHPATPRPEFTSVIRAGVLK
						QDPVRPWGLAPPGDPPPTLLHA
						GFLPPVVRGLCTPGTIRYGPAEL
İ						GLVYPHLQRLGPG\PALPEAFYP
		l		l		PLGLPYPGPAGTRLPRKGD

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23378	53746	A	23513	1319	2211	GSKVYGSLFHINHGNPFNMEVL
23370	33740	ľ.,	23013	1.515		VDSWPEYQMVIIRPQ/ETVRGE
						EKVOGNDLTFHHPTOEMTDDM
		l				DSYTNVYRVFSKDPQKSQEVL
						KNSEIINWKQKLQIQGFQESLG
				ľ		EGIRAAAFSNSVKVEHSRALLF
		1		l .		VTEDILKLYATNKSKLGSWAET
		l				GHPDDELESETPNFKYAQLNVS
		l			i	YSGLVNDNWKLGMNKRSLRYI
		l		İ		KRCLGALPAACMLGPEGVPVS
		1		l		WVTMDPSCEIGMGYSVEKYRR
		l				RGNGTRLIMRCMKYLCQKNIPF
	1	1				YGSVLEENQGVIRKTSALGFLE
		l				ASCQWHQWNCYPQNLVPL
23379	53747	A	23514	234	858	SVPLQESPLPPPPTHQADC*SPC
		l				GAQGGWVTLAVASASS*AHTA
		1				SPLPRSPQAQGRAPSPAPPQIHL
						TTPSSFSSQGT/SHGGLGSTPGA
		1				DDPNTSSS\PPPGGSTPRWSAGD
						SAHSGGEVDGWS*SFP*SVPSY
ł						GT*SPFCPWPSSGQRSFHGIASG
1					1	WEPDGIPDGSSRWEPCGKPQES
1		l				ETGVEGRQHFPKQLCLGSCRA
						VASPRVARA
23380	53748	Α	23515	1	3582	
23381	53749	Α	23516	628	4254	RLPESRAHDVAGADERPWPVH
			l			AGIFKSPAAGAAMSPAAAAAG
						AGERRRPIASVRDGRGRGCGGP
1		1				AGAALLGLSLVGLLLYLVPAA
						AALAWLAVGTTAAWWGLSRE
						PRGSRPLSSFVQKARHRRTLFA
		1				SPPAKSTANGNLLEPRTLLEGP
		1		i		DPAELLLMGSYLGKPGPPQPAP
				ł		APEGQDLRNRPGRRPPARPAPR STPPSPPTHRVHHFYPSLPTPLL
				ì		RPSGRPSPRDRGTLPDRFVITPR
		ı				
23382	53750	ļ	23517	1	392	RRYPIHQTQYSCPGVL KVHMNGLRYALYIMGPALTRG
23382	33730	A	2331/	l'	392	CHOVLIMLCCEVIERILT/VMM
l		1	1	I	ł	KEAYFGEVDIMDDLPT\GNISA
l		1	l		1	NVTGRLYKCGVINHTVDLRRM
						KT/EKWQNNLLPSRQFGFIVLTT
			l	1	1	SAGIMDHEEARRKHTGGKILGF
23383	53751	A	23518	227	352	GROWN DIEE ARREST TOOKIEUT
23363	100/01	l <sub>V</sub>	23310	221	332	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
23384	53752	Α	23519	3	507	SFRAATMVRMNVLADALKSIN
						NAEKRGKRQVLIRPCSKVIVRF
						LTVMMKHGYIGEFEIIDDHRAG
						KIVVNLTGRLNKVGVQWRDLG
						YCNLCLPGSSDSSASASLVAEIT
						\CGVISPRFDVQLKDLEKWQNN
		1				LLPSRQFGFIVLTTSAGIMDHEE
		L				ARRKHTGGKILGFFF
23385	53753	Α	23520	279	653	ARAQRFSSQVDVFYLIQSVIFCR
						LALAEFEIIDDPQSLGKLL*TLT
		1				GRLKQ/CLGVISPRF\DVQLKRP
		1				WKKWHEIIL/PSNPAKFGF\IVLT
İ		1				TSAGLHG/HHEGRQDRKHTGG
		_				KILGILLYRDVIHIFT
23386	53754	A	23521	1	392	KVHMNGLRYALYIMGPALTRG
		1				CHQVLIMLCCEVIERILT/VMM
		1			ł	KEAYFGEVDIMDDLPT\GNISA
	ļ	1				NVTGRLYKCGVINHTVDL/PPY
		1				ENGKWQNNLLPSRQFGFIVLTT
L		L.		L		SAGIMDHEEARRKHTGGKILGF
23387	53755	Α	23522	108	371	LSYIGKFEIIDDHRAGKIVVNLT
						GGLNKCGVISLRFDVQLKDLEK
		1				WQNNLLPSCQFGFIALTTSAGI
22220		١.	20502	252	378	MDHKEAR*KHTAGKILEFFF
23388	53756	A	23523	252	518	HASAFRAATMVRMNVLADAL
23389	23/2/	A	23524	ľ	218	KSINNAEKRGKRQVLIRPCSKVI
	1	1				VRFLTVMMKHGYIGEFEIIDDH
						RAGKIVVNLTGRLNKVGVOWR
1						DLGYCNLCLPGSSDSSASASLV
		1				AEIT\CGVISPRFDVQLKDLEK\
1		1				WQNNLLPSRQFGFIVLTTSAGI
		1				MDHEEARRKHTGGKILGFFF
23390	53758	A	23525	279	709	ARAQRESSQVDVFYLIQSVIFCR
23370	155756	ľ^	23323	1,,	/**	LALAEF*NPLMTHOSLGKFVVN
		1				PHRAGLNKCG\VISPQILTVQLQ
	İ	ı			1	RPWKKWAGLILLP\SROVLGF\I
		1				VLTTSAGI\MDHEEARPKTPRR
		1			1	GKSWGFFFLGDVNYILFYKLKC
		1				LMDSGASTWSF
23391	53759	A	23526	207	564	SMAKEAISINRSLWAGPAKISSP
		Γ.		1		AETE\IESLISLFQKYVGKGGYN
		1			1	CTLSKTEFLSFMNAELAAFTKN
		1			1	OKDPGVLHRMMKKLGTNNDG
		1		1		QLDFSEFLNLIGGLAMACHDSF
	1	1	İ			LKAVPSOKRT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23392	53760	A	23527		407	FREKQSQELSQYGAMGAGQSF NSQFLQHGGPRGPSVPAGMNP TGIGGVMGPSGLSPLAMNPTRA AGMTPLYAGQRLPQHGYPGPP QAQPPAPTGGQENLL*GYSRA AVSARSLPGPFAIPVSHLAGNP TPPMNPH
23393	53761	Α	23528	3	315	
23394	53762	٨	23529		763	DYESTIOPTCSWKP/PVFKPDM IKIEEPOGPALKRCKTVSPAHV LMPSVMEMIAALGPGASPFAPL QPPSVPAPSDYPGQGSSFLGPGT EYGSPIPSSLTSEKSTSLPLQA RRPPAGHLDPTHNPGTPGLTN NLGAPPGPQLHHSNPPASRQS LGQASLGPTGELAFSPATOVMG PPSMSGAGEAPEPALDLLPELT NPDELLSYLGPPDLPTNNNDDL LSLFENN
23395	53763	A	23530		2901	MERRGPGAATARGRARPGGGP SVGLLATGSSLNPSFHGVARIVP GFIRIARPRDGSFAYESVPWQQS ATQPAGSLSVVTTVWGVGNAT QSQVLGNPMGPAGSPSGSSMM PGVAGGSSALTSPQCLQQAFA EGGANKGYVQQGVYSRGGYP GAPGFTTGYAGGPGGLGLPSH AARPSTDFTQAAAAAAVAAAA ATATATATATATATATALQEKQSQE LSQYGAMGAGQSFNSQFLQHG GPRGPSVPAGMNPTGIGGVM
23396	53764	Α	23531	80	412	KPGNGACAGREWCDGGGAAW NWRDPGLPVGDSGVWDRVLLE LLGPRSPRPLDVGGPAAGTPGV LSR/HLPVDCGLGPEALLCSSAP AARRPSLCGGHGRVMRGFRLG CGFRP
23397	53765	A	23532	1	192	RPLDVGGPAAGTPGVLSR/HLP VDCGLGPEALLCSSAPAARRPS LCGGHGRVMRGFRLGCGFRP

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23398	53766	A	23533	1	2094	RPLDVGGPAAGTPGVLSRPCPS TAALAPKPFCAAPRPQPDAPAC AGATGGSCADFDSGVDFVRR SSGLWGPQPPLFPVKNYTEMFQ DPVAFKDVAVNFTOEEWALLD ISQRHLYREWHLDTFWNLTSIG KKWKDQNIEYEYQNPRRNFRS VTEEKVNEIKEDSHCGETFTPV DDDRLNFGKKASPEVKSCDSF VCGEVGLGNSSFNMNIRGDIGH KAYEYQEYGFKPCKCQQPKKA FRYRPSFRTOERDHTGEFWAL LKNDGKKPLFYHSSIQRHMVV
						HSGDOPYKCKFCGKAFHWLSL VLHERTHTGEKPYECQCGKS FSYSATHRIHERTHIGEKPYECQ ECGKAFHSPRSCHRHERSHMGE KAYQCKEGKAFMCPRYVRRH ERTHSRKKLVECKQCGKALSSL TSFQTHIRUMHSGERPYECKTCG KGFYSAKSFQRHEKTHSGEKPY KCKQCGKAFTRSGSFRYHERTH TGEKPYECKQGKAHERTHTG EKPYECKLYGKALSRLISFRRH MRMHTGERPHKCGKAFSS
						PSSYGNEENSH (JEERTYKLAU) GKAFTCFHPFQYHERTHTIGEK PDGCKQCGKAFRSAKYIRIHGR THTGEKPYECKQCGKAFHWVS SFHRHERTHAGEKPYECKHCG KAFTCSIYIRIHERIHTGEKPYQ CKECGKAFIRSSYCKKHERTHTI

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		L			1	
23399	53767	A	23534	48	1317	CGLMDELVHDLTSALEQTSEQ
	1	İ		1		NKLGELWEEMALSPRQQRRQL
				1		RKRRGRKRRSDFTHLAEHTCC
			l	1		YSEASESSLDEATKDCREVAPV
						TNFSDSDDTMVAKRHPALNAI
	1	1				VKSKQHSWHESDSFTENAPCRP
		1		l		LRRRR\KVKRVTSEVAASLOOK
						LKVSDWSYERGCRFKSAKKOR
	1					LSRWKENTPWTSSGHGLCESA
						ENRTFLSKTGRKERMECETDEO
	1					KOGSDENMSECETSSVCSSSDT
	1	İ		1		GLFTNDEGRQGDDEQSDWFYE
		l		ľ		GECVPGFTVPNLLPKWAPDHCS
				ľ		EVERMDSGLDKFSDSTFLLPSR
		1				PAQRGYHTRLNRLPGAAARCL
	l .	l				RKGRRRLVGKETSINTLGTERIS
		1		ļ.		HIISDPRQKDFWLPSAGKRERN
		ı				QFNPLSPLYSLDVLADASHRRC
1						SPAHCSARQANVHWGPPCSRDI
		<u></u>				KQEAETSGHSIFV
23400	53768	Α	23535	225	403	
23401	53769	A	23536	1	403	REQEFLHMAPGEVTITVR\LIRS
		1				LEHR\NFKPVVYHGVNLDQTV
		1		ŀ		KEFIVFLKQDIPLRTNLPPPFRN
1		1		ļ.		YKYDALKIIHQAHKSKTNELVL
	1	1				SLEDDERLLLKEDSTLKAAGIA
						SETEIGIF\REEDYRNYKANPISS
23402	53770	A	23537	2	239	
23403	53771	A	23538	2	214	
23404	53772	Α	23539	1	1341	
23405	53773	A	23540	1	1677	
23406	53774	A	23541	1010	1509	
23407	53775	Α	23542	1	6355	
23408	53776	A	23543	223	10504	VKMPIGSKERPTFFEIFKTRCNK
	1					ADLGPISLNWFEELSSEAPPYNS
						EPAEESEHKNNNYEPNLFKTPQ
						RKPSYNQLASTPIIFKEQGLTLP
						LYQSPVKELDKFKLDLGRNVP
	1			1		NSRHKSLRTVKTKMDQADDVS
	1					CPLLNSCLS\ESPVVLQ\CTHVTP
					l	QRDKSVVCGSLFHTPKFVKGR
1	1	1			1	QTPKHISESLGAEVDPDMSWSS
1	1	1			1	SLATPPTLSSTVLIVRNEEASET
1	1	1	1		I	VFPHDTTANVKSYFSNHDESLK
		1			1	KNDRFIASVT
23409	53777	В	23544	137	709	
23410	53778	A	23545	1	1974	
	1-2.10	٠.	1-22.5	1'	I	

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23411	53779	IA	23546	207	1622	MACCL\SDEVKESKRINAEIEKA
23411	33777	1	25540	207	1022	AAAETRRDARREL/RRLLLLGT
	1	1	i		1	GESGKSTFIK\QMRIIHGAGYSE
		1				EDKRGFTKLVYQNIFTAMQAM
		1			l	IRAM/ETL/KILYKVRSRTRANA
		1				LILIRGGGTLEKVTTFGAFSYVO
		1				LPLKDLFGRDPGIQE/CAYERQP
	İ					RSYQLSDSAKYYLTDVDRIATL
		1				GYLPTOO\DVPAGSRVPTT\GIIE
		1				YPFDLENIIF\RMVDVGG\ORS\E
		1				RRKWIHCFENVTSIMFLVALSE
		1				YDOVLVESDNENRMEESKALF
						RTIITYPWF\QNSSVILFLNKKDL
						LEDKI\LYSHLVDYFPRSSMGPQ
						R\DAKAAREFILKMFVDLNPNS
		1				DKII/YTSHFTCPHRHRRTSRFVF
						PAVKDTILQLNLEEYKPGLSAP
						GPGRRDGDTGQDLPSTEPAAA
		1				RAGGAAESGPGPLPAGGDFFFF
		1			1	IFLTNGFYFTVIRGCTSLPPYTSR
		1				TFSPFVNGKGSLFLALTYGSLFS
		1				KKKKKKKE
23412	53780	Λ	23547	1	401	KKVPGRLSEMSFSLNFTLPANT
23412	33760	ľ^	23347	1'	1	TSSPVT\DCGPSLGLAAGIPLLV
		1		i		ATALLVALLFTLIHRRRSSIEAM
		1				EESDRPCEISEIDDNPKISENPRR
1		1				SPTHEKNTMGAOEAHIYVKTV
1		1				AGSEEPVHDRYRPTIEMERRR
23413	53781	A	23548	584	1071	PCRLPLHLILVEDLSFSLNFTLP
25415	33761	l^	23340	304	1071	ANTVSTAAPIHQAT/PEKTSAEF
i		1				TDCGPSLGLAAGIPLLVATALL
						VALLFTLIHRRRSSIEAMEESDR
						PCEISEIDDNPKISENPRRSPTHE
						KNTMGAQEAHIYVKTVAGSEE
						PVHDRYRPTIEMERRRGLWWL
		1			1	VPRLSLE
23414	53782	A	23549	25	654	LFLOMKDTSPIIFVRKELTIROC
	100	Γ.				TKKVPGRLSEMSFS\LNFTLPAN
		1				TTSSPVTGGKETDCGPSLGLAA
		1				GIPLLVATALLVALLFTLIHRRR
		1				SSIEAMEESDRPCEISEIDDNPKI
	1	1				SENPRRSPTHEKNTMGAQEAHI
		1				Y\VKTVT\KSEEPVHDRYRPTIE
		i				MERRRGLWWLVPRLSLGMMP
		ı				TSNQGSHQTWHLGTGLKTQGF
	1	1	1			VLGEERCOAAS
23415	53783	A	23550	108	413	RRSGSMALWRAYORALAAHP
1-55	1	1	1			WKVQVLTA/GLLGRKAALLAP
	1	1	l			CQ*LYLPTQASWTEGHASCSSL
			l	1		OEATSCPSAFCA*NSKATKOLV
		1	l	1		ERRGLQEHQRGRTLTMVS
	L		L	J	l	Diator delidiron in 12

SEQ ID	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
23416	53784	Α	23551	I	146	
23417	53785	Α	23552	2	442	GREARRSRSMALWRAYQRALA
	1					AHPWKVQVLTA\ASGEEGSTA
l						GPMPVAIPADPGLLD*RTRFLLL
						TPRSHFLPLCLLRMKLKSHKRS
l				l		LMGLGDIISQQLVERRGLQEHQ
l			İ			RGRTLTMVSLGCGFVGPVVGG
						WYKVLDRFIPGTTKVDA
23418	53786	A	23553	3	675	RHSRFGAQGSMALWRAYQ\RT
						LF\ALPWK\VQVLTAGSLMGLG
				l		DIISQQLVERRGLQEHQRGRTL
1						E/TMVSLGCGFVGPVVGGWYK
		i				VLDRFIPGTTKVDALKKMLLD
						QGGFAPSFLGCFLPLVGALNGL
1						SAQDNWGQTTAGELG/RDYPD
l			}			ALITNYYLWPAVQ\LANFYLVP
				l		LHYRV\AVVQCVAVIWNSYLS
l						WKAHRALSLPHSIVSTLQVMQ
					,	LDPWNGSDNPPQKWA
23419	53787	A	23554	512	1227	LQGHLMGIMAASRPLSRFWEW
						GKNIVCVGRNYADHVREMRSA
						/VWLSEPVLFLKPSTGVRAQRA
ļ						RPILMPAYTRNLHHELEL\ALVI
			1			GQ\RCRAVPE\AAAMDYVGG\Y
			1	1		ALCL\DMTAWDVQ\ASARKKG
						LAWDSGE/DAFTASCPVQRVSL
Į.						PKEKIPDPSQ\LKALGFKVNGEL
l			1			RQEGETSSMIF\SIPYIISYVSKIIT
			l			LEEGDIILTGTPKGVGPVKEND
						EIEAGHTRAGQYDHLKVEKAQ
23420	53788	Α	23555	192	436	RTSDHECFREGQICSVCLRERL
						NLNPEEACLLSQAPCMLFVLQY
				İ		LNTVIPYEKKASPPSV\EDLQML
l				l		\TNILFAMKEDNEKVP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23421	53789	A	23556	94	1252	ARMEAPLVSI.DEEFEDLRPSCS EDPEEKPQCFYGSSPHHLEDPSL SELENFSSEIISFSSMEDLVNEF DEKLNVCFNVAKTENLAPV KNQLQIQEEETLQDEEVWDA LITDNYIPSLSEDWRDPNIEALN OKCSDTEIHEKEEEFNEKSEN DSGINEEPLLTADQVIEEIEEMM QNSPDPEEEEVLEEEDGGETSS QADSVLLQEMQALTQTFNNIW SYEGLRHMSGSELTELLDQVEG AIRDFWEELVQQLARRDEE EKEVKNSFFTVLUEVQNKQKIE
						QRELMKKRRKEKGLSLQSSRIE KGNQMP\LKRFSMEGISNILQSG IRQTFGSSGTDKQYLNTVIPYEK K\AYPPSVEDLQMLTNILFA\M KEDNEKVPYFA
23422	53790	A	23557	391	1169	GSFFFFKYFAYSPMPKNSKVV KRELDDDVTESVKDLLSNEDA ADDAFKTSELIVDGQEEKDTDV EGKSEVEDERPAWNSKLQYIL AQVGFSVGLGNVWRFPYLCQK NGGVVPFECEQSSATTYYWYR EALNISSSISESGGLNWKMTICL LAAWVMVCLAMIKGIQSSGKV NTYDRVTMSATVLALQEQWQT FSPIRSGNGHLTGSGAQRTPCRI RRGGSQLRVCDGGKQQWWTV SQSSARAITNTDQKSVQLQDLIE
23423	53791	A	23558	321	568	CLWCGASWPRRHSSPGRPAGD HRGDRP/EGPDTQGPGRPEPSTE PAFNPQSSLDQGGGPAAPAVD DDTRRGHGGVCGDGGAGEA
23424	53792	Α	23559	510	690	LRSPQ\RGVAAPPFLGSPVPFGG FRVQGGMAGQPSAPPALQTDD PYSKAHSVTPKSTPRVT
23425 23426	53793 53794	A	23560 23561	25 84	480 292	VRAVQSLHFHFPWRR/ISCPSES TRSIHVDRLPRRNSASLSLSAFP VSAGALINGEAWVRMGQVPIP KAP

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23427	53795	Α	23562	1138	1924	CRSSATWTPWTTTGCRGDLTK
						TYSLEAYDNWFNCLSMLVATE
		1		l		VCRLLLSPGSAMLRSPATFQVV
				1		KKKHRTRMLEFFIDVARECFNI
						GNFNSMMAIISGMNLSPVARLK
						KTWSKVKTAKFDVLEHHMDPS
						SNFCNYRTALQGATQRSQMAN
						SSREKIVIPVFNLFVKDIYFLPQ
	1	1			ŀ	NP\SNHLPNGHINFKKFWEISRQ
						IHEFMTWTQVECPFEKDKKIP\S
1		1		i		YLLTAPHPTARKLSSSPSFESEG PENHMEKDSWKTLRTTLLNRA
23428	53796	<del> </del> -	23563	3	260	PENHMERDSWRILRIILLNRA
23428	53797	A	23564	1	1386	
23429	53798	A	23565	137	611	MKVNLGDSQNLACVFCRKHD
23430	33/90	1	23303	137	l <sup>011</sup>	DCPNKYGEKKTKEKWNLTVH
						YYCLLMSSGIWQRGKEEEGVY
		1			1	GFLIEDIRKEVNRASKLKCCVC
	1					KKNGAFIGCVAPRCKRSY\HFP
				İ		CGLORECIFOFTGNFASFCWDH
	1					RPVQIITSNNYRESLPWHHLLG
	1				1	NLFEPIPSY
23431	53799	Α	23566	181	486	DPQGEAKGAAPAEGGGGPV*W
						N\GLQGPAGVPGRDGSPGANGI
	1					PGTPGIPGRDGFKGEKGECLRE
		1				SFEESWTPNYKQCSWSSLNYGI
						DLGKIAECTFTKMRS
23432	53800	Α	23567	600	888	QYNGMCSPGPAGVPGR\EGSPG
						ANGIPGTPGIPGRDGFKGEKGE
	1					CLRESFEESWTPNYKQCSWSSL
	1					NYGIDLGKIAGRNCWHILAPVP
		L				NIVSLYFGS
23433	53801	Α	23568	91	1170	ISLHRRPGPTCSMESPASSQPAS
	1					MPQSKGKSKRKKDLRISCMSKP
	1	1				PAPNPTPPRNLDSRTFITIGDRN
						FEVEADDLVTISELGRGAYGVV
	1					EKVRHAQSGTIMAVKRIRATV
						NSQEQK RLLMDLDINMRTVDC
	1					FYTVTFYGALFREGDVWICME
	1					LMDTSLDKFYRKVLDKNMTIP EDILGEIAVSIVRALEHLHSKLS
	1					VIHRDVKPSNVLINKEGHVKM
1			l	1		CDFGISGYLVDSVAKTIN\AGCK VKHGHESLN\PDLNRKGCDVKS
1			1			HVWSLGITMIEMAFLRFPYESW
						GTPFQQLKQVVEHPSPQLPAGR
		1		1		FSPEFVDFTAQCLRKNPAERMS
				1		YLELMEHPFFTLHKTKETDIA\A
						FVKEILGEDS
23434	53802	В	23569	111	193	
23434	23802	ъ_	-5509	1	1.75	

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X-Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23435	53803	A	23570	312	696	RP*SAHPEPGDCPSPPHPPRCGG VAQTAQVPDAPAPV*AG\QHSR SYMEDHLKNKNRLEKEWEALC AYQAEPNSSFVAQREENVPKN RSLAVLTSQPTRNQCLPFLQHG PGDICTPVSISSIVLCHRVT
23436	53804	A	23571		2974	QAAGMGPPLPLILLILLPPR VLPAAPSSVPRGRQLPGRLGCL LEEGLCGASEACVNDGVFGRC QKVPAMDFYRYEVSPVALQRL RVALQKLSGTGFTWQDDYTQY VMDQELADLPKTYLRRPEASSP ARPSKHSVGSERRYSREGGAAL ANALRRHLPFLEALSQAPASDV LARTHTAQDRPPAEGDDRFSES ILTYVAHTSALTYPPGBRTQLH EDLLPRTLGQLQPDELSPKVDS GVDRHHLMAALSAYAA
23437	53805	A	23572		3202	ASRRIPROHGLSAAGQAAGMG PPLPLLLLLLLLPPRVLPAAPSS VPRGRQLPORLGCLEEGLCGA SEACVNDGVFGRCQKVPAMDF YRYEVSPVALQRLRVALQKLS GTGFTWQDDYTQVMDQELA DLPKTYLRRPEASSPARPSKHS VGSERRYSREGGAALANALRR HLPFLEALSQAPASDVLARTHT AQDRPPAEGDDRFSESILTYVA HTSALTYPPGSRTQLREDLLPR TLGQLQPPELSPKVDSG

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	endon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23438	53806	Α	23573	I	1459	MPLTSQDVDQKTAAWVPTNRD
						STVPSDPWDCKRRPWPLLVLRP
	1		Î			LAPGTKAPPGGQNGRGRQKGS
	1		i			PCSRRLLLEPAGPLGPASVRCIR
						TTTPOLLIITTSLRFPSLEAAVGY
	1					GSNTCIFANACQMVFIMITWPY
1				i		LOCIPYSVLLKDLEMRNLRELE
				1		DLIIEAVYTDIIQGKLDQRNQLL
1		-	ł	1		EVDFCIGRDIRKKDINNIVKTLH
				l		EWCDGCEA\ALLGIEQQVLRAN
1				l		QYKENHNR/TLQQQVEAEDVYI
1						LPQSGNDCRGLRFVSSLIGGVR
		1				CVHLKHKGDAVVWEAIEEGEA
					1	LPGKYRLKGGRSVWAAAVRA
						VCMDQVTNIKKTLKATASSSA
1			ŀ		ł	QEMEQOLAERECPPHAEQROP
ŀ			i		l	TKKMSKVKGLVSSSPLGPAGA
1			l	l.		AGTHQAWVRWGGDTKGPFPPL
				l		STCSEFOTCPSPHORLPTLLVLF
1					ŀ	
						QKNCYSPSPTPSFPSCSLQTQGL
						HQCHPKTGSDTAQLPSRRFLSL
						CKGLVSLPVFLLLPRHFVRLVIK
23439	53807	Α	23574	2	396	
23440	53808	Α	23575	1	1014	
23441	53809	Α	23576	110	212	
23442	53810	Α	23577	1	747	
23443	53811	Α	23578	323	495	GRAGKWHTQPE*SLRKAM*MF
1		1		1	İ	QILMMKMIVALARATSGTEAV
1		1	l	l		AGRSWASGLRRSGRV
23444	53812	A	23579	26	229	
23445	53813	Α	23580	I	574	
23446	53814	A	23581	889	1189	
23447	53815	A	23582	170	625	GAAVALVRPNGGTAASHVVCR
23447	1550.5	ľ.	20002	1	1020	EMRPLRPLOSRRCSRRRTGISN
ľ		į.				MRALENDFFNSPPRKTVRFGGT
1	1	1	l	I		
1		1		l	1	VTEVLLKYKKGETNDFELLKN
1	1	1		1		QLLDPDIKDD\QIINW\LLEFRSS
		I		1		VMYLTK\DFE\QLISIILRLP\WL
		_				NRSQ\TVVEEYLAFLGNLVSA
23448	53816	Α	23583	166	455	RWKPSGTWPRNPMERPGH*ES
		l		l		VPKPGD*AKKSGGET\RGRKEG
1		1		I		NEHRETFSQIPDCHLNKKSQTG
		1	ĺ			VKPCKCSVCGKVFLRHSFLDRH
		1		1		MRAHAGHKRS
23449	53817	A	23584	3	3706	
	1-2011	<u> </u>	1	<u> </u>		l

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X-Unknown, *-Stop endun, /-possible nucleotide deletion, \-possible nucleotide insertion)
23450	53818	A	23585	11	1970	MVTDRKAAAGTHGEARPCWG
23430	33010	^	23363	1,	1970	SGSRVHTYRGAGERAMRVHRI
				i		MPREDSERVREADGGKHEORS
						FGDLHPTA APAPA ASGCGTQP
				İ		RVKQPGEWQHREAPQQRQPP\
		İ			1	RPHGSHQLVTVRYERRPGGKE
					1	RARLRPGLGCPSGRTPQGRAP
	1				ŀ	GTALPTAAAAAIFPSAALTSGQ
	1	l				RNASLPKFYFRGGTWGTSQSE
		l				LCGKLSNQFLGEPLVPRIRLILS
		l				CGTCSGCSSPRKDPGHPEAGN
		l			1	PSRDFFQLPSEERDSVAFEDVA
		1				VNFTQEEWALLSPSQKNLYRD
		1				VTLETFRNLASVGIQWKDQDII
		l				NLYQNLGIKLRSLVERLCGRKI
	ŀ					GNEHRETFSQIPDCHLNKKSQT
	ŀ	ı				GVKPCKCSVCGKVFLRHSFLD
		İ				HMRAHAGHKRSECGGEWRET
	ŀ					RKQKQHGKASISPSSGARRTV
						PTSMRPYE\TKVCGKA\FIS\PIY
		ı				QIHQRTPTEKRSYKCREIVRAF
						VSSFFRKHGKMHTGEKRYECK
	l					YCGKPIDYPSLFQIHVRTHTGE
	İ					KPYKCKQFGKAFISAGYLRTH
	İ			l		IRSHALEKSHQCQECGKKLS\C
		l				SSLHRHERTHSGGKLYECQKC
		1		l		AKVFRCPTSLQAHERAHTGER
		1		1		YECNKCGKTFNYPSCFRRHKK
		1		j		HSGEKPYECTRCGKAFGWCSS
	ļ	Ļ				RRHEMTHTGEKPFDCKQWVK
23451	53819	Α	23586	86	390	LGSGDLPWEINPLSSCSLLREK
		l		l		PPTTSGPQT\TSPRNISPISNPEL
						THAGNLATGPRNARSPGFLLSI
		1		ŀ		VPSVWDPTENRTVQLTWQPLP
22152	50000	١.	22507	ļ	500	ELLELWPKVL
23452	53820	A	23587	1	502	MLLTQSLFGGLFTRTHMKFGA
		1				VTQIRGPPLGDKSPVLLLFALE
	1	1				QRRHVLSMDPKLRCWSRTGK
		1			1	AFPWCLIIAEMPDYSPTFQRCQ
		1				TTQGRLPWSFTLSSKSRFSGEG
		1				ARACYKCQKSDHQAKECPQPO
						IPPKPCPICAGP\TGNRTVQLTW
		<b>⊢</b>				QPLPEPLELWPKAL
23453	53821	В	23588	1	663	
	53822	Α	23589	233	496	
		IA.	23590	2	378	ERQRRHVLSMDPKLRCWSRTC
23454 23455	53823	1.	i .			
	55823	ľ				KAAFPWCLIIAEMPDYSPTFQR
	153823					CQTTQGRLPWSFTLSSKSRFSG
	53823					

SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23456	53824	Ā	23591	3	452	VRPRRDACLGPSPLAASPAFLG
				ľ		KGQVPQPLISLCPDPLFPHPNLIS
		l				LRPNPLCPHPDLVSLCPDPFPAF
						LEAHKNFOTPEPOOPGIPPEPPP
						PGACYKC/HEI*PPGOGMPAAO
1		l				DSS*AMS/LSVRDPTGNQTVQL
						TWOPLPEPLELWPKAL
23457	53825	A	23592	1352	1540	TWQTELEDESWITE
23458	53826	A	23593	177	383	EAMEPWKGVPLGRVNTEQIAP
23430	33020	ľ.	23373	1'''	500	FSEAEGPREIH*GSRKVRKAAO
1						GLNEGPRELFWOPLPEPLELWP
		l				KAL
23459	53827	A	23594	1478	1744	ICAE .
23460	53828	A	23595	23	178	SCLEVCDEOGPEK\TRORALRG
23400	33020	ľ	23030	123	1.70	VSSVTEDTLNICRLCWOPLPEPL
		ļ				ELWPKAL
23461	53829	A	23596	1	282	EEWIKAE
23462	53830	A	23597	34	241	SGDLPWEINPPSSCSLLREKDPP
23402	33030	ľ`	1337,	1	1	MTSGPQT\TSPRNISPISNPDPTG
		l				NRTVOLTWOPLPEPLELLPKAL
23463	53831	A	23598	121	286	AASFYSSPTSLTIPOPLSPFNLGV
23403	33031	ľ``	23370	1.2.	200	TLOSLPSLNFNSF\LFWOPLPEPL
		1				ELWPKAL
23464	53832	A	23599	144	488	GYEIWCRDSDOGTFPWEINPPS
23404	33032	ľ.	23377	1	1.00	SCSLLREKDPPTTSGPOT\TRPR
				1		NISPISNLVSGLFLLSSPPSLTIPO
		l				PLSPFNLGATLQSLPSLNFNSFH
		1		1		FLVETKETRFIRGAKIPAPVTD
23465	53833	A	23600	471	677	TEVERNETIA INGILIATI II TIE
23466	53834	В	23601	67	555	
23467	53835	Ā	23602	1	1440	
23468	53836	Ā	23603	860	1209	IPGNCRDSSGVGIKERETNAGS
	1	1.		1		OHMH/IKNSSVHGYPRLCVEDE
		l			1	DAYKKOFSOYIKNSVTPDMME
		l		1		EMYKKAHAAVPENPVYEKKPK
	1	l				KEVKKKRWNRSKMSLAOKKD
		1		1		WVAQKKASFLRA
	<u> </u>	L	L	1	L	THE PROPERTY OF THE PARTY OF TH

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23469	53837	A	23604	1	966	MGFVKVVKNKAYFKRYQVKF
1	ŀ	1				RRRREGKTDYYARKRLVIODK
1		1				NKYNTPKYRMIVRVTNRD\IICC
1						IANA\RIEG\DMIVCAP\YAHELP
		1				KIWV*RVGLTNYAAA\YCTGLL
		1				AGPAGFFHRFGMDQIYE\GQV\E
		1				LTGDEYNVESIDGOPGCLHPAY
		l	1			WDARPLPRTYPLAIKVFGCP*R
		l				GA\VDGGLVLFP\HSTKTIPPGY
		l				DS\ESK\EFNAES/IHRKPHPWAQ
		l				NVARLHAPTLMEEDEDA/YTK
		l				KQFSQYIKNSVTPDMMEEMV/Y
		l				RKAHAAIRENP\VYEKKPKKEV\
		l				KRKRWNRPKMSLAQKKDRVA
		l				PKKASFLQSSRSGLLESLTPAIFF
23470	53838	A	23605	1	644	ANSSPVNPVVFFDVSIGGOEVG
		Γ.		ľ		RMKIELFADVVPKTAENFRQFC
		l				TGEF/RVSFRCV*LLCLLORGVL
		1				GLO*GYR*TTCWP*MIAGDSDR
				1		RKDGVPIGYKG\STFH\RVIKDF
		l				MIQGGDFVNGDGTGVASIYRG
		l			i	PFADENFKLRHSAPGLLSMANS
		l				GPSTNGCQFFITCSKCDWLDGK
		1				HVVFGKIIDGLLVMRKIENVPT
		l				GPNNKPKLPVVISQCGEM
23471	53839	A	23606	638	1391	CPAYKGAHGIQCVYCTLPPPLL
						QTTRRVRRNPTLPTSASGSEPW
		l				RLANSSPVNPVVFFDVSIGGOE
						VGRMKIELFADVVPNTAENFR
1		l				OFCTGEFRKDGVPIGYKGSTFH
		l				RVIKDFMIQGGDFVNGDGTGV
		l		ł		ASIYRGPFADENFKLRHSAPGL
1		l				LSMLTWPGKLTAGIFGFHQANS
		l		İ		GPSTNGCOFFITCSKCDWLDGK
		l				HVVFGKIIDGLLVMRKIE\GMPC
1						IFQPTWTHFLPVITTACSALGSM
1						LAAERGTKN
23472	53840	A	23607	2	591	ARAEFGTSRVGAMAVANSSPV
		l				NP\VVFFDVSIGGQEVGRMKIEL
		ı		1		FADVVPKTAENFROFCTGEFRK
		l		I		DGVPIGYKGSTF\HRVIKGF\MI\
		1		I	Ì	QGGDFV\NGDGTGSRPVFTRGP
		l		l		ILQMKIFKL\RHSAP\GLLSMAN
		1				SGP\STNGC\QFFITC\SKCDW/V
		1		I		WDGKHVVFGK\IIDGLLVMRKI
	l	1				E\NVPTGPNNKP\KLPVVISQCG
23473	53841	A	23608	6	424	19499
	1000.1	1		I*		L

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide togation of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
23474	53842	A	23609	3	470	DAWVSGRLTELPSSCFFPPIFRO
				Ī		LLPVMPEPSKSAPAPKKGSKKA
						VTKAOKKDGKKRKRSRKESYS
i						VYVYKVLKQVH\PDTGISSKAM
						GIMNSF\VNDIFE\RIAGEASRLA
l						HYNKRSTITSREIQTAVRLLLPG
1						\ELCKHAV\SEGTNA\VTKYTSS
23475	53843	A	23610	93	42 I	
23476	53844	A	2361I	2	1173	DHAAPRRSSRASVATVTRAGR
						CPTMSVRVARVAWVRGLGASY
l		1				RRGASSFPVPPPGAQGVAELLR
l		l				DATGAEEEAPWAATERRMPGQ
		l				CSVLLFPGQGSQVVGMGRGLL
1					1	NYPRVRELYAAARRVLGYDLL
						ELSLHGPQETLDRTSQVQSPGT
		1				GIPRRVRAHVLHLSPQVIENC/V
						VS/AGLAFSPSFCALIFPNALL/F
						WPGLYAVKIRAEAMQEASEAV
						PSGMLSVLGQPQSKFNFACLEA
Į.						REHCKSLGPVNGQICKGGFLGH
						CSAETSFLPQALRFLQKNSSKF
						HFRRTRMLPVSGAFHTRLMEP
						AVEPLTQALKAVDIKKPLVSVY
					İ	SNVHAHRYRHPGHIHKLLAQQ
					İ	LVSPVKWEQTMHAIYERKKGR
		<u> </u>		105	105	GFPQTFEVGPGRQLGAILKSCN
23477	53845	A	23612	187	425	Maria de la porte de la composición dela composición de la composición dela composición de la composición de la composición de la composición dela composición dela composición de la composición de la composición de la composición de la composició
23478	53846	A	23613	1	1258	MKAYLLLGAAPPLPWFPGQAV
		l				FRPCPPPAGLWRTAQLVELEHG PVPREDLIPCSRPFSISFLGPERV
		l				DGLAFPRIQALGIAAARGMEPK GAPGKSDSGNSRSAGGRVLGC
		l				OGAALEVECSAGRGGPVRLAV
		i				GCVLLRAPERGWRRGEGAARG
						IPAGRSSESLGALSPQVGVRSDI
						GINGNIWHPGVOTDFSSLKRSO
						GLYAVKIRAEAMQEASEAVPS
						GMLSVLGQPQSKFNFACLEARE
						HCKSLGIENPVCEVSNYLFPDC
			1			RVISGHQEALRFLQKNSSKFHF
						RRTRMLPVSGAFHTRLMEPAV\
		1	l	1		EPLTQALKAVDIKKPLVSVYSN
			i			VHAHRIQGIPGHIHKLL\AQQLV
			l	1		SPVKWEQTMHAIYERKKGRGF
			ĺ			POTFEVGPGRQLGAILKSCNMQ
					1	AWKSYSAVDVLQTLEHVDLDP
	L				l	IN WAS I SAVDVLQ I CERVOLUP

SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
			sequence		
53847	Α	23614	2	1050	GRCPTMSVRVARVAWVRGLG
					ASYRRGASSFPVPPPGAQGVAE
					LLRDATGAEEEAPWAATERRM
i					PGQCSVLLFPGQGSQVVGMGR
					GLLNYPRVRELYAAARRVLGY
					DLLELSLHGPQETLDRTVHCQP
					AIFVASLAAVEKLHHLOPSVIE
					NCVAAAGFSVGEFAALVFAGA
					MEFAEG\LRFLQKNSSKFHFRR
					T\RMLPVSGAFHTRLMEPAVEP
			1		LERKLLKAVDIKKPLVSVYSOR
					PTRIRLOAFPGNIOOSCLAOOLG
		-	1	Į.	LPK*KWEQ\TMHAIYERKKG/R
					RGFPQTFEVGP\GRQLG\AILKS
					CNMOAWKSYS\AVDVLOTLEH
					V\DLDPQEPPKMTGRGLKCNDP
ŀ					LCPPEERL
53949	Δ	23615	600	905	OAPASGSPWLSGGPOOVEDAG
33040	ľ	23013	000	J~~	AGYGFAPGOPPPPRTOPRSAC
					SRRAAG/TPVPRAPPAPSAG\PK
l	1				ARAPRSLGCGSLGRFSTGVRPD
ł					KCIFPPETENAAGCF
53849	A	23616	390	722	QAPASGSPWLSGGPQQVEDAG
			-		AGYGFAPGQPPPPPRTQPRSAC
					SRRAAG/TPVPRAPPAPSAG\PR
					AAAPRSLGCGSLGRFSTGVRPD
					KCIFPPETEMSYLLFRNTVSRCR
					c
53850	Α	23617	1	673	MGQVKGHKLCLDFTLLSPDGG
					AAEGRCLELGWOAKEVEETIE
1					GMLLRLEEFCSLADLIRSDTSQI
			i		LEENIPVLKA\KLTK\MRGIYAK
					VDRIKAFVKMVGHHVAFLEAN
					VLQAERDHGAFPQALRRWLGS
					AGLPSFRNDPTAPLRQVFEASP
1					DPRLARTVTIQTQQPVRVSGPL
					SGQRKEKSLQKGAQVFQTNVP
					ETLLEGAAISPAPACGGPLTSGD
					GFRHVRK
53851	Α	23618	1	651	
53852	Α	23619	195	497	DQFLENTKKCDGLNQLSFDPA
	l				NNNKPLEFGRARGPLVTEGLIE
	l				NGGESSKKRRRTNSSSADNSRT
					LNVDSTAMTLPMSDPTAWATA
1	1		1		M\NNFGMAPVRIVGH
	53847  53848  53850	53847 A  53849 A  53850 A	of peptide sequence         bod in USSN 809/540,217           53847         A         23614           53848         A         23615           53849         A         23616           53850         A         23617           53851         A         23618	of peptide sequence         bod         in ISSN 109/840,217         location of first sequence           53847         A         23614         2           53848         A         23615         600           53849         A         23616         390           53850         A         23617         1           53851         A         23618         1	In USSN   location of first codon for last amino acid sequence

SEQ ID		Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last aminu acid	*=Stop codon,/=possible nucleotide
	sequence	l	09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23485	53853	Α	23620	241	1823	NDDRAKFTVEDHTMQRDFRW
						LWVYEIGYAADNSRTLNVDST
						AMTLPMSDPTAWATAMNNLG
		ĺ				MAPLGIAGQPILPDFDPALGMM
						TGIPPITPMMPGLGIVPPPIPPDM
		1				PVVKEIIHCKSCTLFPPNPNLPPP
		l				ATRERPPGCKTVFVGGLPENGT
		ĺ				EQIIVEVFEQCGEHAIRKSKKNF
						CHIRFAEEYMVDKALYLSGNR
		l				ELKKFLNKEDNYFRKINLTALY
		l				KTVLEERCEKRRGGYRIRLGSS
						TDKKDTGRLHVDFAQARDDLY
		l			į.	EWECKQRMLAREERHRRRMEE
						ERLRPPSPPPVVHYSDHECSIVA
		l				EKLKDDSKFSEAVQTLLTWIER
					l	GEVNRRSANNFYSMIQSANSH
		l				VRRLVNEKAAHEKDMEEAKEK
						FKQALSGILIQFEQIVAVYHSAS
		1				KQKAWDHFTKAQRKNISVWC
		l				KQAEEIRNIHNDE\LMGIRREEE
		l				MEMSDDEIEEMTETKETEESEL
		l				KKLEGIVDGSVEVSAEALRKLL
		i				NEYERQVKEEQKVGYRIWGTI
						HNKGEGKKEEFAEELEKSSSER
23486	53854	Α	2362 I	996	I512	RPVHLASPAKEQQGTRKGRRA
		l				PARLPVPTTGRGLEKGPESQAP
		i			l	PSFRQATGRFSRSGSSGGPTPSK
		l				NSRDCSHTPQLPTPRGNPYVFPS
			ĺ			PGHPEPSRPVEMPAPG\EPRTRQ
		1			i	LCRERTRKACPPKPRPPLGLPG
						DPTGPVTHHAPPVSPTGASGQE
			l		1	RRAEPGAVSYAHASATK
23487	53855	Α	23622	293	1140	PRGILHQDKNLVVINK\PYGLPV
						HGWPWG\QLCITDVLPILAKML
		1				HGHKAEPLHLCHRLDKETTGV
		1		ŀ		MVLAWDKDMAHQVQELFRTR
		1				QVVKKYWAITVHVPMPSAGVV
		l				DIPIVEKEAQGQQQHHKMTLSP
		1				SYRMDDGKMVKVRRSRNAQV
		l				AVTQYQVLSSTLSSALVELQPIT
						GIKHQLRVHLSFGLDCPILGDH
		l				KYSDWNRLAPQKLSVGTLKKL
		l				GLEQSKARYIPLHLHARQLILPA
						LGSGKEELNLVCKLPRFFVHSL
		1	1			HRLRLEMPNEDQNENNEAKCL

SEO ID	ISEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
		1	i	sequence		
23488	53856	A	23623	1	1328	MAVDLGSHTPGSSGVTAATFR
23488	22820	<u>۱</u> ^	23023	1'	1328	
1	l	l				GRRLVNTREESVVLGAGGEVL
		1				SGVPGTWDALGPGASFGIKGM
		1				FPKGTVIMSTSPMCEGPVPRRK
		1				PCASSVGQQNSRSRKKAGRKTP
		1				TLAGRHVPVKIKKEAIQQKPLP
						ASSFSPALTRSGAIFGPCPALTS
		1		ļ		APVLEYPLQKVVENEAFKVFN
		1				NRDEESKRQKQAEFRMLTSAIR
			1			GPAGPRGHSSTRKPPSNPPPPGA
		l				CFKCGNEGHRSRQCPNPGTTSQ
	1	1	i			EAELIALTRALTLAAGQQINIYS
		l				NSHYVFHIVHSHSSIWKERGILT
	1	l				AKNTPAINGSLINKLLKAARLP
		l				QKVAIIHCRGHQTPDNPILAGN
		l				VLADQVAKQVALQPVQGQFLS
İ		1		1		LSLFSPLYSSEEKEDFRAQNLQ
ĺ				1		KQGPW/LQSQGARLLVPAACAS
	1					KVAEAGRKPAGRHVPLKIQKE
	1	1				AIWSSPFSVPMATRAAASTSPT
23489	53857	A	23624	475	622	IHSKVLFSASSSSC*NTNNSWFR
		1				ASIKKNPEKONKNMOSAIPOHG
		1				LWKI
23490	53858	Α	23625	17	992	GRGEVLRGRGRGQLDAVLSRE
1	1					HCGGSRQSEQPWSRRSSSSSSR
		1				RRLRSAAEPAMALSMPLNGLK
		l				EEDKEPLIELFVKAGSDGESIGN
i						CPFSOG\LFMILWLKGV\VFSVT
İ			i			TVALKRKPADCQNLVPGTPPP\
						LILFSSEVKTDVNKIEEFSEEVL
		İ				CPPKYLKLSPK\HPRSNTAGMDI
						FA\KFSAY\IKNSRPEANEALER
1	1	l				GLLKTLQKLDEY\LNS\PLP\DEI
1	ŀ	l				DENSMEDIKFSTRKFLDGNEMT
1	1	1		1		LADCNLLPKLHIVKVVAKKYR
1					1	NFDIPKEMTGIWRYLTNAYSRD
1	1	1				EFTNTCPSDKEVEI\AYSDVAKR
1		ı				LHOVKSRLLKEVSFMSSP
1	I	ı	I.	I	1	LHQVKSKLLKEVSFMSSP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
	1			sequence		
23491	53859	A	23626	11	1128	MALASGEGLPVALSHGRRGNG
		i .		ľ		OERVPYLKLRDTCAYRVAWRC
						NRLARAKVKRTSYRSGYGNGF
		1	l			EPGVEPDCVVLLALGVENDLV
				1		RISSKAEKYSLYEFSFPLRVSRE
İ		1	l			CEEECRKEPWFNPGOKPSVLIT
		l	1			SGQILGQEEQKTLLFPGLLSLVF
1		1				OFNVENSDLRCLSLAENLEISAL
1		1				LSLSCSSVIRLILINLEVTISPPITI
1		1	1			
1			1			LKHAPPLTSLEASSSSSGIEKKIH NNFROGPFPSSATRSTFLMNHT
1		l				SFDSYVVVSNIVPRLNLYPOVL
		i				
		1				DVLQSLHKIHQLASNAKWTIQG
		l	i			PAKPDLGLELGLHRYPDWEKE
1		1				ERKKKRKRSVSPEKSITTHNKP
1		1				TSSGIFY/RPRRCGCDLGRPPAP
		Ļ.				GPHPSSPPHSPPQPSRCPPV
23492	53860	Α	23627	3	453	SLGKDIRPRSARAACKGVGLWS
		1		l		GCFGKMAGSGVRQVTSTASTF
1		ĺ		1		VKPIFSRDMNEAKRRVRELYRA
		1				WYREVPNTGER*RLVRPAVLK
		_				HSYVANRSF
23493	53861	Α	23628	3	695	PYASRRKFRVDPRVRAWSGCF
		ı		1		DKMAGSGVRQATS\TASTFVKA
1		l		1		HFSVGDMNE\AKREGARVLTAP
1		ł		1		WYR\EVPNT\VRVHQFQLDI\TV
		1		1		KMGRDKVREMFMKNAHV\TDP
1		1				RVVDL\LVIKGK\IE\LEETIKVL
		1				KQ\RTHVMRFFH\ETEAPRPKDF
		l				LSKF\YV\GHDPLKSFSGKDAR
		ł				WILFLGAQIKLTIQWSLCDRIPL
l		l		1		MNQFSAASFCLSKWGPWVLHI
		L_				TCSFPFSWKCGVCCKLD
23494	53862	Α	23629	82	441	SLPINPVPTACTLQPAGLLRPPG
				1		SDVRTPMLPPSAPSQPSYPTSRP
1		1				PLWPTWPAGPVFRRSQPLAQPN
l		l				LPRSYHQQAPGGQTPVALGLE\
		ł				CPKATNQPFGTLQRDPKSPAKV
		1				PPFEEFSL
23495	53863	Α	23630	3	504	DRSSSSTMSTGGDFGNPLRKFK
		ł				LVFLGEQSVGKTSLITRFMYDS
	i	1				FDNTYQATIGIDFLSKTMYLED
1		l				R/TNVNSFQQTTKWIDDVRTER
		1				GSDVIIMLVGNKTDLADKRQVS
1		1				IEEGERKAKELNVMFIETS/AKA
				1		GYNVKQLFRRVAAALPGIMPS
		1		1		VLFQIIPSQALTRYLC
	<u> </u>				L	. 5. 4 54

SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino seid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	delction, \=possible nucleotide insertion)
				sequence		
23496	53864	A	23631	416	1239	GGGRSSSSTMSTGGDFGNPLRK
						FKLVFLGEQSS\GKTSLITRFMY
	l					DSFDNTYQATIGIDFLSKTMYL
		1				EDRTIRLQLWDTAGQERFRSLIP
						SYI\RDSAAAVVVYDITNVNSFO
						ANYKSGIDDVRTERG\SDVI\IM
						LVGNKTDLADKRQVSIEEGERK
	1					AK\ELNV\ILMETSAKAG\YNVK
						QLLSTCKQPALPGMESTQDRSR
						EDMIDIKLEKPQE\QPVSEGGLF
						LTNLPCHLQPFFQKLTGFWPPY
1	ŀ					SFIDCSVNIGLMPFPFSNNVLRFI
		1				IAACPAWR
23497	53865	A	23632	1	493	PKPATAPPPPSL/RARFLCPORRP
23497	33863	<u>۱</u> ^	23032	['	493	ERAR/PRARGASPRPEKTAGAR
i						KRRGSCERPHPGG/RPPAPG*I
	1					SSRPTLGFFWILNFFFLKKLGRI
						KDVPWODSTKEHLSVCRGLRG
1						TWSVPSASDPLHPAWRRTCFRV
1				1		PPTTGAREWWLHLPPPALORPP
		1		1		DSROCHRIPVC
23498	53866	A	23633	1	560	MESIWAGLWTAODTTVLADDF
23498	22800	I <sup>A</sup>	23633	'	300	EFILWTLKRRPERAR/PRARGAS
		l	1			PRPEKTAGARKRRRGSCERPHP
		1				GGLRRLPGARDQRAPPRSDAG
		l				RSPGRLRRLPALLCAGPLSPGRE
		l				PRRWEPRAGVARPAGQQWGFF
						WPGNQQGASICSLILWFLLECG
		1				SRSTGGPCSGTELGHFSRERPTV
		1	·			LLSRLGTVKRP
23499	53867	A	23634	32	327	LTORLFGGLFTRTHMKFGALTR
23499	33807	<u> ^</u>	23034	32	321	IG\DLPWE\INPLSS\CSLLPEKDP
1			i			PMTSGPQTDQPKEHLTNFKSAA
		l				RRSSRTSSHRNLLHVLEIWOLD
						QGMPAARDF
23500	53868	Ā	23635	2	424	LTPFSDSAHLHPGEINSPVAHTK
23300	33800	^	23033	<u> </u>	424	PVGGLFTRMRVTFGAKTODRS
1		l	1			TP\QETSPLSSPS\PVTSGPQTNQ
1		i				PKEHLTNFKLGKQSFHSSHASL
1		l		1		ATLOSSSLATLOSPCPSNSNSFS
1						SLVETRRRPGAVARLQSQLSGR
						LRGGRIA
22501	62000	ļ	22626	31	246	
23501	53869	A	23636	Pi	240	TALLLTQSLFGGLFTRTHMKFG AVTRIG\DLPWGINPLSSCSLLR
	1		1	I		
	1		1	I		EKDPLTISGPQTHQPKEHLTNF
L		L	L	L	L	KSGPH

	SEQ ID NO:		SEQ ID NO:			Amino acid sequence ( X=Unknown,
	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	coden for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
23502	53870	Α	23637	125	464	TAVFLTQSLFGGLFTRTRMKFG
						TMTRI/RGDLPWEINPLSSCSLL
						HEKDPPTTSGPQTNQPKKHLTN
						FKSRKVERTKGLLKTHLTKLSH
						QLKKAWTILLPLSLLRIQACPR
						NAT
23503	5387I	Α	23638	1064	1304	TAMLLTQSLFGGLFTQMCMKF
						GAVTRIG\DLPWEINPLSSCSLL
						HEKDPPTTSGPQTDRPKEHLINF
						KSGACYTCRKSGH
23504	53872	Α	23639	539	747	TAMLLTQSLFGGLLTRTRMKF
		ł				GALTRIG\DLPWEINPLSSCSLL
				1		HEKDPHTPSRKIHIRPQGNHESQ
						KGL
23505	53873	Α	23640	709	950	TALLLTQSLFGGLFTRTHMKFG
						AVTQIG\DLPWENQSPVLLCSLL
1 1						REKDPPTTSGPQTDQPKEHLTN
						FKSGKWSFHSSPAF
23506	53874	Α	23641	1	1257	MGKKVLGIASQLLGAVRCEPD
						GRRVLPEDWPECLSLVTHGTW
						FTRQGDKNIQMADNSFSDGVPS
						DFVEAAKNASKTEKLTDQVMQ
						NPGVLAALQERLDNAPHTPSSY
						IETLPKAVKRRINALKQIQVRC
						AHIEAKFYEEVHDLERKYAAL
					ļ.	YQPLFDERREFITGDVEPTSDM
			1	l	1	ESEWHRENEEEEKLAGDRKNK
						VVITEKAAATAEEPNPKGIPEF
						WFTIFRNVDMLSELVQEYDEPI
		l				LKYLQDIKFHFQPNDCFANSVL
		1		ŀ		TKTCKMKSEPDKADPFSFAGPE
		1				IMDCDGCTIDWKKGKNVTVKT
				l		IKKKQKRKGPGTVRTITKQVSN
						ESFFNFFNPLKASGDGESLDKD
		İ				SEF/ASDFEIGHFFRER\PQAVLY
						FAGEALKDDDNFDEGEEEEDEL
						EGDEEGEDEDEVEINPNKEPSQ
	53875	Α	23642	132	1277	
23508	53876	Α	23643	1	515	RVRHRSVSVGSTHASAHAYGE
						GSLRIIPVTRSPLVTMNHIVQTF
						SPVNSGQP\PNYEMLKEEQEVA
				l		MLGAPHNPAPPTST\VIH\IRSET
				ĺ		SV\PDHVV\WSLFNT\LFMNTCC
				l		LGFIAFAYS\VKSRDRKMVGDV
						TGAQAYASTAKCLNIWALILGI
				L		FMTI\LLIIIPVLVVQAQR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	coden for last amine acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23509	53877	A	23644	243	643	NHTVQT\FFSPVNSGQPPQRMR
						CIKEEHEVAVAGG\PHNPAPPTS
	l	1			ŀ	T\VIHIRSETSVP\DHVVWSRVQ
		1				TPSSMNPC\C\LGFIAIRLTP*SLR
						DRKMVGDVT\GAQA\YAST\AK
						CLN\IWAHDSGASVMTILLIVIPS
23510	53878	A	23645	3	1498	RCNFLRSSRIRVHPTPAASTMPP
					l	KFDPNEIKVVYL\RCTGGEVGA
	İ	1				TSALAPKIGPLGLSPKKVGDDIA
	l	1				KAMGDWKGLRITVKLTIQNRQ
						AQIEVVPSASAMIIKALKEPPRD
	l				ľ	RKKQKNIKHSGNITFDENGNIC
	İ					STDMR/HRSLSRELSGTTKEILG
						TAQSVGCNVDGRHPHDIIDDIN
	İ					SENDKQPRKSKMRTSFSPQIPG
						HEHFHYLLLLQHVHCQLALTL
						RARKEGSSTHHQKLSTYYFKEG
	l					NSKEPKKIQRCRNTELRKEEGG
	l	ļ			l	GGSGKAPGSEIKLESPACSGASL
						GEMTAWTMGARGLDKRGSFF
		1				KGPLPNTCGHFWEMVWEQKSR
		1			İ	GVVMLNRVMEKGSVVEFVTEL
	l	1				FFYVALPKMFPVHYIVILLASKL
						EEVTFNFPPPPDLCTPFLQNYLR
	i	i				VAFQEVNSGCTGKTLLVRPYIT
	i					TEDVCQICVEKFKVGDPEEYSL
					ŀ	FLFVDETWQQLAEDTYPQKIKA
		1				ELHSRPQPHIFHFVYKRIKNDPY
						GIIFQNGEEDLTTS
23511	53879	Α	23646	361	918	GCTGSRGTVPLSALGPQRIGPP
						GSCLQIKSWVNDICQRQPG\DW
			İ			KGP*GFTVKLTHSRTGQAPDCT
		1				RVP/VLPSCP*FIQSPSREPPKRPE
		1				KKPEKTFKHQLGIFTFLNEICST
		1	i			LLRQMR\HRSLSPEKLSWNPT*
		1	l		i	RDSWGIAQFSGAVNVDG\RH\P
		I				HDI\IDDIQQWVLWKCPRQLST
		1		_		KENIFNKGII
23512	53880	Α	23647	1	642	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23513	53881	A	23648	894	2438	VETKRIQPQPPDEDGDHSDKED
23313	33001	l'`	250.0	0,,		EOPOVVVLKKGDLSVEEVMKI
ŀ		ı				KAEIKAAKADEEPTPADGRIIY\
		1				RKPV\KHPSDEKYSGLTASSKK
		1				KKPNEDEVYQDSVKKNSQKLII
İ		1				NSSLLSFONEDENDTEKSMENL
		1				TGKIRKLQLRIVDPTFLPKLSNK
		l				GKDPVAVARLETGKGNKGAFR
ŀ						VLRDISRWATLCTSPQHSSSEA
						KNIIPYQATQFQNEVHKWYKK
		1	l			KGGLKISKRKALFFLTQDIQKK
		1				LOKLALGPNTPTPDILKIASLVF
1		1	l			YNODQEEKERAQQNERQKEKL
1		1				YWLSYKSASPFQVALRILSQET
		1				THKDELPTDLPNNPEVQAPGIP
		1				GNTIMAMPVAIQLKDPSSYPCR
1		1				IIKVPEIASWIHRTQVKLWKSPE
		1	ł			TSELEPTPSVIEYTREALEDLKF
ł		1				LFKOKDNSSAKTPGRAYAASPR
						DWTSIEAELHIFYOWDPNLKG
		1				MFKPASLLAKVKODFPDIHKES
		l				PIFRAIFSNITLMGIAPICVTGKTI
i				}		
1	ĺ	i				KKRTRQHHTTPKSNLREQASQ GSLNWEGN
23514	53882	A	23649	200	1727	RTATSLDSVLYKSDYMTWKYR
23314	33002	ľ.	23047	200	1''-'	VSFFYILSCRLLKK/QCTDNEHV
						YMELSKKLYKYCPKEWKKEAS
	1				İ	KVRQYEVTWVVSKRGKDYILK
						HIPNMHKDOFALTASEAHLKYI
						KEAVRLDDVAVHYYRLYKDK
l			i			REIEASLTLGLTMRGIQIFONLD
		ŀ				EEKQLLYDFPWTNVGKLVFVG
l						KKFEILPDGLPSARKLIYYTGCP
		l				MRSRHLLQLLSNSHRLYMNLQ
		l				PVLRHIRKLEENEEKKOYRESYI
		1				SDNLDLDMDQLEKRSRASGSS
	l					AGSMKHKRLSRHSTASHSSSHT
						SGIEADTKPRDTGPEDSYSSSAI
						HRKLKTCSSMTSHGSSHTSGVE
I		1	l			SGGKDRLEEDLQDDEIEMLVD
1		1	l			DPRDLEOMNEESLEVSPDMCIY
I		1	1			ITEDMLMSRKLNGHSGLIVKEI
		1				GSSTSSSSETVVKLRGOSTDSLP
1		1	l			OTICRKPKTSTDRHSLSLDDIRL
I		1	l			YOKDFLRIAGLCODTAOSYTFG
1		1	1			CGHELDEEGLYCNSCLAOOCIN
		1	l			IQDAFPVKRTSKYFSLDLTHDE
23515	53883	A	23650	182	423	RLSIMASSLNEDPEGSRITYVKG
23313	22003	l^	23030	102	723	DLFACPKT\DSLAHCISEDCRM
			l			GAGIAVLFKKKFGGVQELLNQ
	1	İ	1			HYKEKGLRTSQLYG
	L		L		l	III VEVOLVIOČE IO

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23516	53884	Α	23651	1	249	
23517	53885	Α	23652	435	494	
23518	53886	Α	23653	163	578	RVTWLKKHLNSGNSDSVCMAS SLNEDPEGSRITYVKGS\LFEGP KTDFLAPCISEDCRMGAGIT\VL FKKKP\GGVQELLNQQKKS\GE V\AVLKRRWGRLYIFTWITKRK
						RAFAQSPLYENLTEEFRGQLKS HLSEEWSH
23519	53887	A	23654	1	2238	HL3EE W3H
23520	53888	A	23655	26	1941	RDAGTKWQRLGHLSCHCRKQA AREIINMHFQAFWLCLGLLFISI NAEFMDDDVETEDFEENSEEID VNESELSSEIIYKTPQPIGEVYFA ETFDSGRLAGWVLSKAK\KVD
						MDEEISIY DORWEIEEL KENQV PGDRGLVLKSRAKHHAIS AVLA KPFIFADKPLIVQVEVNFQDGID CGGAYIKLLADTDOLILENFYO KTSYIIMFGPDKCGEDYKLHFIF RHKHFKTGVFEEKHAKPPDVD LKKFFTDRKTHLYTLYMNPDD TFEVLVDQTVNKGSLLEDVV PPIKPPKEIEDPNDKKPEEWDER AKIPDSAVKPEDWDESEPAQI EDSSVVKPAGVLDDEPKFIPDP NAEKPDDWNEDTDGEWEAIPQ LINPACRIGCGEWKPPMIDNEY YKGVWIPPLVDNPNYQGIWSPR KIPNPDYFEDDHIPILTSTSALG
23521	53889	A	23656	297	398	ADHWAADGWRWKIMIANANK PGYLKQLMAAAEGHPWLWLI) LVTAGVPIALITSICWPTKVKDI RHSDLEYIIPDICIPQTKGVLEQE EKVEKAALEKPMDLEEEKKQN DGEMLEKEESSEPFEKSEEIEII EGQEESNQSNKSGSEDEMKEA DESTGSGDGPIKSLEKRRYRKD
23522	53890	A	23657	286	430	TAGARCSSRTSSPGSCFNCOKS
23522	53890	A	23658	405	524	GHWAR\NARSPGFLLSRVPSVN WIHI ARMSVACLPGLFFICLRGDFKQ
		1				NLTAYCWKCYQL*RSLGTHS
23524	53892	Α	23659	ı	657	
23525	53893	A	23660	22	334	ALKHFCLCSLIFSVTTMKFLAV LVLLGVSIFLVSAQNP\TTAAPA DTVSSLLVLLMMKPLDAETTA AATTATTAAPTTATTAASTTAR KDIPVLPKWVGDLPNR

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nuelcotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequen <del>ce</del>	deletion, \=possible nucleotide insertion)
23526	53894	A	23661	I	270	
23527	53895	Α	23662	1	284	
23528	53896	Α	23663	111	281	
23529	53897	Α	23664	229	363	
23530	53898	A	23665	250	586	PVRHGAFFQDKSSDKKVQTKG
				[ ·		KRGAKGKQA\EVAN\QETKEDL
Į.						PAENGETKTEESP\ASDEAGEKE
	1					COCLINNHIPCLISGPCLPSCTIO
		1				RNIFINYFVNASFLVALETFLKK
23531	53899	A	23666	3	478	APAVGLSSSGGGRSGSGKAARF
23331	33677	l^	23000	ľ	770	REGCRRAAARMQQAPGTRLPR
ŀ		1				RHPARAFPAATMPKRKVSSAE
	i	1				GAAKEEPKRRSARLST\KLPAK
		1				V\EAKPKKAAAKDKSSDKKV\Q
		1				TKGKRGAKGKQA\EVANQETK\
	ł	l				EDLP\AENGETERLEEESSLLIEA
						GEKEKPKS
23532	53900	A	23667	1	252	GERERINS
23533	53901	A	23668	328	414	
23534	53902	A	23669	1	3156	
23535	53903	A	23670	1887	2036	
23536	53904	A	23671	3	212	
23537	53905	A	23672	22	539	YEGIVMKPLIIVNPADCIGC\RT
23337	35705	ľ.,	23072	[ T		CE\VACVVAHPSEQELNADVFL
1	i					PRLKVQPIWTAVSAAR*CCPFSS
						ENAP\CVGA\CPVGA\LTMGEQV
1					i	VQTNSARCIGCQSCVSACPFGM
}						ITIQSLPGDTRQQIVKCDLCEQR
1						EEGPACVESCPTOALOLLTERK
						LRRVRQQRIVVSGENPL
23538	53906	A	23673	135	302	PIHMKTROPRDOFSADCVFLOS
				1		TCPVPC*RNRL*TPGGYWRTRR
						DYLRCDRSVHV
23539	53907	A	23674	175	794	LSTHSQPIADAADPPPHLHTGEL
				1		LCSAHOADFAVSAEPARPANN
				1		NAATTAPVHAVAITPPAYPDCF
						RTLPKSLSHODSTORLYGIARL
						LLRISTSGPPPSRLKVORLDSISA
						PV/MCHOCENAPCVGACP/VGA
						LTMGEQVVQTNSARC/IGCQSC
						VSACPFGMITI/OSLPGDTROOI
l						VKCDLCGALRDCVAVLRORRN
						SRVVLDACTRA
23540	53908	Α	23675	2	434	LISLPFPPDSLAHRPPSSAPTEPQ
	1	Γ.		Γ	1	GLFTVAAPTPGAPSPPATLAHL
l	1	1	1	1		LPTPAMYSLLETELKNPVGTPT
I			1			QAAGTGGPAAPG/GRKMALEN
l						PKMHNSEISKRLGADWKLLTD
1	1		1	1		AEKRPFIDEAKRLRAVHMKEYP
1				1		DYKYRPRRKTKTL
23541	53909	A	23676	3	264	DIKIN MICHEL
1-7741	123707	LC.	22070	Ľ	,	

SEQ ID	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23542	53910	A	23677	1	2377	MDDTFLNYKYKNKCGPKHIKK
						KEEEEEEDNNSNDSKNKDKTL1
		1	İ			TGSSGSAKIFLMRGGRRQLLLP
		l	i		l	TLSLCLLSVQRSPGSHESGPPQS
		1				MNPRLSPQGKYLKNDFLDVIG
		1				MCDIACHLVVELLVGNNGNFL
ļ		1				AHAFAHVEVVAQQNQRENAA
			Ì			RPNGGSVMIGQGASLRTCQSRV
		1				LRVARGGPSPNPGESEQGLFTV
						AAPAPGAPSPPATLAHLLPAPA
						MYSLLETELKNPVGTPTQAAGT
			l			GGPAAPGGAGKSSANAAGGAN
		1				SGGGSSGGASGGGGGTD\QDR
					1	VKRP\MNAFMVWSRGQRRKM
		l				APENPKMHNSEISKRLGADWK
		1				LLTDAEKRPFIDEAKRLRAVHM
1						KEYPDYKYRPRRKTKTLLKKD
1		1				KYSLPSGLLPPGAAAAAAAAA
1				1	1	AAAAAASSPVGVGQRLDTYTH
						VNGWANGAYSLVQEQLGYAQ
į.						PPSMSSPPPPPALPPMHRYDMA
						GLQYSPMMPPGAQSYMNVAA
İ			İ			AAAAASGYGGMAPSATAAAA
						AAYGQQPATAAAAAAAAAAAM
			ŀ			SLGPMGSVVKSEPSSPPPAIASH
				l	1	SQRACLGDLRDMISITGRENIFV
		l			1	VGDPAACLLPIEMEETFLPWPL
		1		1		GFEDRLHLEGPRPVQEIPSFLRK
		1	l			VPNCGQWLLQTPSVEPADPWE
		1	1			NPQAFPPSLQQEQPCLSLIVGIF
		1		ĺ		VEAFGGRGRWGSKGKKPKRYR
		1				TVEWMRIHQPSICCLQETHLTR
23543	53911	A	23678	1	399	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
23544	53912	A	23679	13	1216	LGNKRAGLRRADGEAPSALPY
						CYFLQKDAEAPWLSKPAYDSA
	1					ECRHHAAEALRVAWCLEAASL
						SHRPGPRSGLSVSSLRPTRKMA
	1					TTFLAHEKIWFDKFKYDDAERR
						FYEOMNGPVAGASROENGA\T
						VILRDIARARENIHKSLAGSSGP\
						GAS\SGTSGDHGELVVRIASLEV
						ENQSLRGVVQELQQAISKLEAR
						LNVLGKELRLAHRGTAPOTOH
						VSPMR\QVEPP\AKKPATPAEDD
			l			EDDDID\LVGRDNE\EEDKEAA
						OL\REER\LR\HYAEKKAOESLP
	i					LGGQVPPSLLDVK\PWDDESDM
						AQLEACVRSIQ/LGTGRSWGAS
			l			OMVAVGYG\IRKLOIOCVVEDD
	İ					KVGTDLLEEEITKFEEARARSV
						DIASFSTRSEALKCVYVAHVRE
						GPCTRSKDLRPAKKKKKK
23545	53913	Α	23680	1	2066	MAAALRAPTQVSAASSRPPLPS
						PPETEGRVKGFLQPGPHCPAQC
						LCSGDDGSNEDLEQKREVIFPR
						PHEAPVAEHRQQTVGLRELVAI
						PKRKEETTAIVQVFVAFEDVAI
						YFSQEEWELLDEMQRLLYRDV
						MLENFAVMASLGCWCGAVDE
	l					GTPSAESVSVEELSQGRTPKAD
			İ			TSTDKSHPCEICTPVLRDILQMI
	1					ELHASPCGQKLYLGGASRDFW
	İ					MSSNLHQLQKLDNGEKLFKVD
				ı		GDQASFMMNCRFHVSGKPFTF
1	l			1		GEVGRDFSATSGLLQHQVTPTI
			l			ERPHSRIRHLRVPTGRKPLKYT
						ESRKSFREKSVFIQHQRADSGE
						RPYKCSECGKSFSQSSGFLRHR
	ŀ		1			KAHGRTRTHECSECG\KSFSRK
	l	1				THLTQHQRVHTGERPYDCSEC
						GKSFRQVSVLIQHQRVHTGERP
	[	l	İ	l		YECSECGKSFSHSTNLYRHRSA
	İ	l				HTSTRPYECSECGKSFSHSTNLF
		l				RHWRVHTGVRPYECSECGKAF
		l				SCNIYLIHHQRFHTGERPYVCSE
	1	l				CGKSFGQKSVLIQHQRVHTGER
	1	l				PYECSECGKVFSQSSGLFRHRR
		l				AHTKTKPYECSECEKSFSCKTD
		l		1		LIRHQTVHTGERPYECSVCGKS
		l		I		FIRKTHLIRHQTVHTNERPYEC
				l		DECGKSYSQSSALLQHRRVHT
		l				GERPYECRECGKSFTRKNHLIQ
	1	ı		l		HKTVHTGERPYECSECGKSFSQ

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
NO.	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
23546	53914	Α	23681	Ī	468	MAAQVGAVRVVRAVAAHEEP
	ì	1				DKEGKEKPHAGVSPRGVKRQR
						RSSSGGSQEKRGRPSQEPPLAPP
l						HRRRSQPPTSWAAAVNDM/PAP
1						TVPGPVEPLLLPPPPPPSLAPAG
	1					PADDAPLPAPSTSGLFTFSPLTA
!	1					SAAGPKHKGTQGAAQAPSPPRP
						RW
23547	53915	Α	23682	950	1326	RPESQRANGVDSGPNLKTVPQP
	1	i				DTRKGSVLKWISKRGKPLAVEI
l			l			EESHCL\CLPLRTECLGIKP\IVH
					1	LFSCTRPVIVPSLELNYDVDSIA
l						HMFVADLLLMITLPSYDIPFYC
		<u> </u>				LVFQNLLVLEFQYL
23548	53916	A	23683	271	918	NQGLRDLGLCRICLVNKMFTSS
i						ILGKGHRHSLVSIKQEHSALRK
	l	1			1	AAG\PLPLKVGYC\QGFSPCDSL
	1	l				KYG\SWDEKDLTVPQPDTHKGS
1		l		1		VLRWISKRGKPLAVEIEEG\HCL
		I				PELPPGELECPG/ILKHGLYHWS
		l				SSEMGE\NRPMVG\ARHVYSNA
		ı				ALLFFTALRIFGQRRNINLGLRA
	1	i				HPG\IVPPLELNYDIDSSAHM/FF
20540	53917	١.	23684	,	942	LADLLLIITLPSYYIPFC
23549	53917	A	23685	346	1753	TWIPLSRTGRRFPVFSCSSSRYV
23330	33916	ľ	25005	370	1755	AAISFHKRWKEGRCFGRTRGE
1			İ			MGWEGRSRSLKELNTRNDWNP
		l		ì		GERELDFWRKITRFKEAASWTL
		1				EALEEAGSTARRRVWLEDGRR
i		1	1	Ì		TCEILSVLCTSSLSSNCASLDCV
		1	İ	Į.		TYKALSTTRLYDSVYWNKKPE
	İ		1	ļ		RSSSPATEOSWTENDFDKLREE
		1		l		GFRRI\NYSELKEEARTHGKEVI
				l		NLEKK\LDERLTRITNAEKSLKD
		ı	1			LMELKTMARELRDECTSFSSQF
		1				DPLEERVSVMEDQMNEMK\QE
		ı				EKFREKRIKRN/EKQSLQEIWDY
1		ı				VKRPNLHLIGVPESDRENGTKL
		İ		1		ENTLQEIIQENFPNLARQANIQI
				1		QEIQRTPQRYSSRGATPRHIIVR
1					1	FTKVEMKEKMLRAAREKGRVT
1		1		1		HKGKPIRLTADLLAETLQARRE
		1		1		WGPIFNILKEKNFQPRIAYPAKL
				1		SFISEGEIQYFADKQMLRDFVTT
		1		1		RPALKELLKEALNMERNNWYQ
	1	1		1		PLQKHAKW
-						
23551	53919	A	23686	73	327	GALQPTAALWEPLSGLAKAAA
23551	53919	A	23686	73	327	RSLSLQGGVEGDARAGTGAAR
23551	53919	A	23686	73	327	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23552	53920	A	23687	322	402	DQEPTNSGHIGNLEQCLGHGYQ R*LFP
23553	53921	В	23688	545	3063	
23554	53922	Α	23689		579	MKHEKRDRCAELTAILLNEQAS PMCTIVLGYTGTSTITLPRTSCH FAPRLSNSQUESSYQVGHKPES YCVRNWWVLGLTDFKNEAAD PHNSGQQLASPSGSRTGAAGG\ VPASPAPCARTPOPLGGRWAD APGTGGGIRRGGSGCTGTHGG GGRLRHGGLQSGGLPRGKAAK ARREJESSAGYSNSSSKCSII
23555	53923	A	23690	399	592	KCIHFPGPTPKFVCGVIVVTAVP /TFGM*TVYV**LPFSILAQNVG *RDDCDPGWKLNLFYAKKKK
23556	53924	А	23691	3	268	DRV/RPLRKNFQRNDQTAAFTV HENPLLCSYRCWYPGKQEQKK IKPQRGGKNRAEKLETLKSRAP LLFQRNAVPHQQRNKAGWRM TLTS
23557	53925	А	23692		484	CSFFLVSMVFTFWHDFAAAGTS CSFPCLVLSFRSSFRAGLVVTKS LSICLSVKYYFISSSLMKLSLAGY EIVG*KFFSLRMLNIGPHSLLAC RVSAERSAVSLMGFSLRVTRPF SLAALNIFSFISTLVNLTIMCLG VALLEKYLCGILCVS

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion. \=possible nucleotide insertion)
23558	53926	A	23693	182	2076	MQEWLRLTARKTELSRDSRSC GQLKTSLLISLLVESTOGRI+SF LVWTOTHSGGDGWRVAKAW RFKYHDTQLGGEKEEPQGHHA SDCRWRAQPHCRTLSAAVVNV AGFTVTFELGCLTLMKMREKD HHLCDNDIVLEPPDTAFPLDCL LSCHISTVHGLLRWNAIQYLAS SESSHPERSEGTPDRNOHPAL SLASFNGAQODRRTVQSLGEKP GLPEEMAAFMAGTEKEQNELG IEFFQLIPASKRTHSENALFSHS GDDTECSRSQQCSPPSYSRGPP RTFILATOPLCSKEAGTSPHEQ PTRTEKNRGSHLYHKSPCDHAS PGMSVTMSGMQQALPRQLTIL THYPPAFRSIKFIRMILKTEMLH THARSQEALLKDSSKTMLHSTL PSCVRKSNYHHRERKIKTEMAL FAKHRSNGAMKVEVAPLITI SSDPLAEFILLPVTVTLCSASLEW LAPEGGMLSUNDERSENSEW LTLPEGHSGLLIPLSQVKRV
						PVLVGVIDLDYQDEIDYYSSMK ERSSPATEQSWMENDFDELRE EGFRRSNYSELREEIQTKGKEV ENFEKNLEECITRITNTEKCLKE LMELKTKVRELREECRSLRSRC DQLEERVSAMEDEMNEMK
23559	53927	Α	23694	933	1118	RVERRRLQTIKLLRATGGNSNQ RQRS*KL*KKFRRMYN*NNQY REVLKGADGAESQGSRTT
23560	53928	A	23695	2	250	WCHTKLDGLSLFGESVALEEA GGQTALWLMRFLSRKANYVD KITTREVGHRE*LMSSRLHSR* WAASSVCSSSCVLLPSSHS
23561	53929	A	23696	1017	1230	
23562	53930	В	23697	62	1069	
23563	53931	A	23698	216	582	ILVPTLFWLGEFLLRDPLLV+W ASI*G*PD/AFSLAALNIFSFISTL/ RESDNYVSWSCSSRGVSL*RSL YFLNLNVGLPCQIGEVLLDNIL QSVFQLGSILPVTFGYTNQMQI WFHFIHFIFHH
23564	53932	A	23699	434	681	TLACLARLGKFSWIISCRVFSNL VPFSLSLSGTPIRRRFGLFT*SHI SWRLCSFLFILFSLNFPSRFISFV SSSITDTLSSS

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23565	53933	A	23700	290	1231	RSFRAIGEVGSIRGWGGGYAAO
1		1		I		LQSPSRLPVLGGGKKGGSRGV
		1				GOGSSRAEVGLGSQRERSGTCG
1						AEAAAAAAQLIAGTGSSLPYQG
1		1				PGONOTAIAPTCAPLSPCKAAA
		1				VIMGNIFGNLLKSLIGKKEMRIL
		1				MVGLDAAGKTTILYKLKLGEIV
İ		1				TTIPTIGFNVETVEYKNISFTVW
		1				GCGVARDK\IRPL\WRH\YFON\T
		ļ				QGFDILWSDSN\DR\ERVNEA\R
		1				EEL\MKNAGAETSLRDAVLPFL/
		İ				FGNKTDLRNAMNAAG\QDKL
		l				G\LHSLRHRNWYI\QATCATRG
						NGLYEGLHWLANSVPKQEVES
						RTALTKHPNRP
23566	53934	С	23701	164	208	
23567	53935	С	23702	1	381	
23568	53936	С	23703	43	276	
23569	53937	С	23704	1	462	
23570	53938	С	23705	38	121	
23571	53939	A	23706	77	202	QKNYTQRPLPLHGLLELEVTCR CWSWQPLLEPLELRP\RLSD
23572	53940	Α	23707	123	209	
23573	53941	С	23708	1	411	
23574	53942	Α	23709	1400	1809	SSNPLSRPPLPTPVTPCPLLRLSL
		1				LRACCCCLFLPLRFFSCPLLCFL
			ĺ			LCFFFSVCSPAHKNFKMPKPQW
		1				SGIPTGPPPSGSCFKCQKSGHW
ì		1				A\RNACSPGFLLSHVPSVWDPT
		1				GNQTVQLAWQPLLEPLEL*P\R
		١.		_		LSD
23575	53943	В	23710	76	358	
23576	53944	В	23711	1	606	
23577	53945	С	23712	272	388	
23578	53946	Α	23713	1	259	PGVRGAQGGPSIPRQCEESAIGP
						KFAWIISSFHWKQGMFNMEAP
		1				SSLFFVNMCAVKKQTTWGRPD
		_				RGFIWQPLPEPLEL*P\RLSD
23579	53947	Α	23714	684	917	GDFGLRQWGFLNIQSCHLQTGT
		1				I*LPLFLIEYPLFLSPA*LPWPELP
		1				TLCCIGVVREGISVLCQFSKGM
		<u> </u>				LPVFAHSV
23580	53948	A	23715	618	683	
23581	53949	A	23716	306	710	DRVSLLTPRLGVQWPDPGSLQP
		1				RPPGFKRFYCLSLPSS*DYRHEP
		I		L	<u></u>	LLPAYFFPYRIIGWSRL
23582	53950	Α	23717	2	88	

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first eodon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop coden, /=possible nucleotide deletion, \:=possible nucleotide insertion)
23583	53951	A	23718	972	1967	FPPGRGROPGRAGCSRGRARA CPKPVDHVKRPMNAFMVWSR AQRRKMAQENP;QMHNSEISKR LGAEWKLLTESEKRPPIDEAKR ILGAEWKLLTESEKRPPIDEAKR ILKAMHMKEHPDVKYRPRKFK ILLKKDKFAFPVPYGLGGVAD AEHPALKAGAGLHAGAGGCH PSELLANPEKAAAAAAAAA VFFPQSAAAAAAAAAAA VFFPQSAAAAAAAAAAAA VFFPQSAAAAAAAAAAAA VFFPQSAAAAAAAAAAAA VFFPQSAAAAAAAAAAAA VFFPQSAAAAAAAAAAAA VFFPQSAAAAAAAAAAAA VFFPQSAAAAAAAAAAAA VFFPQSAAAAAAAAAAAA VFFPQSAAAAAAAAAAAA VFFPQSAAAAAAAAAAAA VFFPQSAAAAAAAAAAAAA VFFPQSAAAAAAAAAAAAA VFFPQSAAAAAAAAAAAAAA VFFPQSAAAAAAAAAAAAAA VFFPQSAAAAAAAAAAAAAAAA VFFPQSAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
23584	53952	A	23719	2	144	RPQRRP/LPPPTGSPLKHPGGAS GAAGAPIGEPQHPGAVVSVRA WSLCS
23585	53953	A	23720	219	897	PGRAGGTLGLOSRRDOTREKG VEETPHRLHARTDTTAPGCCGS PIGAPAAPEAPPGCFEGPRWVE GVKKASPPLPAELIGWLVNFPL ARPVAPPSLSOPPRRPPEGVCG SWPGRSEGERGKAKNTAESTPR QNLHRKYTIIVTHYGRNRKGQ NISFSLVCEFIPSLADWTTVAL LTWTSPKDNSLTEESWLREGN VHKGMDEYVRAQSFLILMTLE NSGGSVVSK

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
ł	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23586	53954	Α	23721	937	2730	ТИРНСИДНҮҮНИННЕНСИНСИ
					ŀ	нининсининунининской
	ĺ		ŀ			нүннинисинсинининсин
				ì	ŀ	ННҮНННҮНННСНДННҮНННН
	1		l			<b>НСИДИНУНУНИНИСИДИНУН</b>
	1					ннинсидинүнүнинсидин
ļ						Y НИНИНСИDНИЧНИНИНСИD
	l			i		ннүнининсноннүнүнинс
						НННКҮНННННСННСННННН/Р
1			l	i		нинининисниннунининс
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						ннинүүнинсинсининүн
						ннинсирннунинсинсини
						НҮНИННҮНДНИҮНИННЕН
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1		i				ннисинсунинуниннунси
1		1				ННҮННННҮНРҮНҮНОНННН
Į.		l				ҮНРННҮНННҮҮНСННННОҮН
						НРННСНННННҮНОННҮҮРІТІІ
						TTTITTITTIIITVTTTTFTTTTNT
		l				TAPTTTPVTAITTTTTTTTTTTTT
		1				TPITITTITNITTTTVTTTVIITTTF
		l				ТТТІНРҮННННННРУТІТТІТ
		l	İ			TATHTTTTSITSITTHTTTIVTTH
1		1				ТТТААТНИНҮНРҮҮННҮННН
1		1				ННННТНЅҮҮНҮНННННННСГ
						НҮНИНҮНИН
23587	53955	A	23722	2348	2571	
23588	53956	A	23723	4919	5142	
23589	53957	A	23724	2	370	
23590	53958	A_	23725	588	701	
23591	53959	A	23726	295	521	
23592	53960	A	23727	2837	3061	
23593	53961	A	23728	272	436	
23594	53962	A	23729	239	514	TOP DOCUMENT OF STREET
23595	53963	Α	23730	105	367	TGRPPELLLHLRSLSPPLL/CSSP/ TPPPQLPPPPVSLASGYPLLHPL
		1	1		l	ARYPACPLPPRRARTEORGGEO
						PGVGASPLPPRRAKTEQRGGEQ
22.506	52064		22721	999	1146	POVGASPLPIPGKWPGTEAT
23596	53964	A	23731	1084	1145	
23597	53965	A	23132	1084	1893	l

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \mpossible nucleotide insertion)
				sequence		
23598	53966	A	23733	1605	2296	HSIHSSVDHPENRIYRRKIEELS
23376	33700	*	23,33	1005		KRFTAIRKTKGDGNCFYRAL\AI
1	1					PYLESLLGKSREIFKFKERVLOT
				1		PNDLLAAGFEEHKFRNFFNALR
					· ·	AGRAOFYSVVELVEKDGSVSSL
						LKVFNDP\GASDHIVQFLR/LCF
			1			TSAFIRNRADFFRHFIDEEMDIK
			ł			DFCTHEVEPMATECDHIOMTA
						LLQALSIALQVEYVD\GWIPALT
						OHVFLEAATPSVYMLYKTSOY
	l					NILYAADKH
23599	53967	A	23734	529	1830	AGYGDRTMLARKSIIPEEYVLA
23399	33707	<u>۱</u> ^	23/34	329	1050	RIAAENLRKPRIRDRLPKARFIA
			1			KSGACNLAHKNIREOGRFLODI
		1				FTTLVDLKWRHTLVIFTMSFLC
			- 3			SWLLFAIMWWLVAFAHGDIYA
						YMEKSGMEKSGLESTVCVTNV
		l				
		1				RSFTSAFLFSIEVQVTIGFGGRM
						MTEECPLAITVLILQNIVGLIINA
						VMLGCIFMKTAQAHRRAETLIF
			l			SRHAVIAVRNGKLCFMFRVGD
						LRKSMIISASVRIQVVKKTTTPE
						GEVVPIHQLDIPVDNPIESNNIFL
		1		}		VAPLIICHVIDKRSPLYDISATDL
		l				ANQDLEVIVILEGVVETTGITTQ
		l				ARTSYIAEEIQWGHRFVSIVTEE
		l				EGVYSVDYSKFGNTVKVAAPR
		l				CSARELDEKPLPSLFRPSQKSEL
		l	l			SHQNS\LRKRNS\HRRNNFHRRN
		l				N\SIRRNN\SSLMVPKVQFMTPE
		_				GNQNTSES
23600	53968	A	23735	968	1243	author is the state of the stat
23601	53969	Α	23736	541	1010	SVTSCLKVTSVCLTSAAASCRH
						RRSVEEYLVS*KLPM/WSRRRK
	l	1	1	I		AAALTES/V/CIVFWLASWMMS
	1	1		I		RWLHYPDWVTIIAQQVLMNVV
	l	l	l	l		SVMQADCRMNLLEIFLRWLMR
1	1	1		l		KIFHNILRHFSQLLNVPITRPFS
		_				WILRPF
23602	53970	Α	23737	766	1910	LLTARLTWVSARLMSYVLLMC
	l	1		l		*LFPRLLSCLTTPPHCSFSICFVI
		I	l	l	l	CSRTLILKGSSLMYVFCLPNTAI
		I		l		GGSSRVDRCRAATSRGSSRVDL
		I	l	l	l	TL*KYSLVGPSVPLVSSV**SDW
		1		l		PLKLLLMTMMS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23603	53971	IA	23738	581	1434	I KLPMGVAVVKLLHLPKVIIVFW
						LASWMMSRWLHYPDWVTIIAQ
						OVLMNVVSVMOADCRMNLLEI
		ł				FLRWLMRRASDICSSIHGD*EPL
		1				L/YAVFGKQNTYIRLEPFKINVL
i		1				EQITKHIEKLQCGGVVKQLSRR
ļ.	1	l			i	GNNOHISSTYDINRADTOVRRA
		1				VNNYDIIVMSNSFNGQSEHQV
1						WIDRKAGECLLHEYEDLVPIRD
1					i	TLRLFPGGRYLPRAKHVAPSEP
		ı			l	DPEQDEQKLRFCKRHLYGKOP
1		1				RSPVERVWDHGPTRIVGLIISLG
		i	ľ	1		PRSHSYRRSDNQTGTTVPLVSS
23604	53972	A	23739	515	774	LFPRLLSCLTTPPHCSFSICFVIC
23004	337/2	l^	23139	1313	l''*	SRTLILKGSSLMYVFCLPNTAIS
		1		1		GTTVPLVSSV*LLVWDHGPTRI
		1		i		VGLIIRLGPRSHSYRRS
23605	53973	A	23740	322	763	CRMC*LFPRLLSCLTTPPHCSFSI
23605	339/3	A	23740	322	163	CFVICSRTLILKGSSLMYVFCLP
1						NTAIVMALSPRGWRSKFGMPV
1		1				EDTSLSTPAAPMVDSLIARVGV
[		1				MARGNAITLPVCGRDVKSTLE
l		1				VLRGDSVEKTSRVWSGNERDO
		ı				ELLTEDALDDL
20.606	53974	+.	23741	11	127	MKLMETLNOCINAGHEMTKAI
23606	539/4	A	23/41	11	127	AIAQFNDDSPEARKITRRWRIG
		1		1		EAADLVGVSSQAIRDAEK*RRL
		1				SPLHSLMMTARKRGK*PGAGE
23607	53975	A	23742	9	437	WLGLGNWKERLGTOTRMIWS
23007	33913	l^	23/42	ľ	437	OKTARRLCRTFSGIPAYSMCI/CI
						GMAERMWCDRNRERHTVSSSG
		1		1		GNRLPNPDWAPGPPRIVGLNIS
						LGPTVPTRIVGLIISLGPRSHSYR
						RSDNQTGTTVPTRIVGLILVLGP
						WSHSYRRSDY
23608	53976	١.	23743	766	767	LLTARLTWVSARLMSYVLLMC
23608	53976	Α	23/43	/66	/6/	
l		1		1		*LFPRLLSCLTTPPHCSFSICFVI
1					1	CSRTLILKGSSLMYVFCLPNTAI
	1	1				GGSSRVDRCRAATSRGSSRVDL
				1		TL*KYSLVGPSVPLVSSV**SDW
		_				PLKLLLMTMMS
23609	53977	A	23744	573	1085	VCHQILTHLPALLSGARMMWD
						QLWRGRP*RRCQQPQCGG*WH
1		1			1	CFWATAHHSPLDSTLNHLHCLP
						RGP
23610	53978	Α	23745	2	379	WLTRSSCSWRLPPRRKKKGAL
		1			1	V*PVWLPPGGVAVLLQLGCCV
		1		1	1	LQDVSCKRLCYNRKGQSEIVL
		1		1	1	MS*ITMWKLFPRPPLLPFSTCIC
		1		1	1	*KTCGKRGRSK*KVDILVLSIRF
		1		1		SAIKLEPLDCHENAWL

SEQ ID	ISEO ID NO.	Inter-	SEQ ID NO:	Nucleotide	Nucleotide location of loca	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		丄			L	
23611	53979	Α	23746	575	576	TGIDSDQLNGRDWRRALSVSPA
						PDSGFPRCNGDEQAGI*LLGTG
				1		WAEKCHRLAFPPAGSTAGR
23612	53980	Α	23747	1	723	MKLMETLNQCINAGHEMTKAI
		i .		1		AIAQFNDDSPEARKITRRWRIG
		1				EAADLVGVSSQAIRDAEKAGR
		1		1		LPHPDMEIRGRVEQRVGYTIEQI
		1	1	1		NHMRDVFGTRLRRAEDVFPPVI
		1				GVAAHKGGVYKTSVSVHLAQ
		1				DLALKGLRVCSWKERPPGNSL
		1				NVSRMVWDHDPTRVVGLIIGL
		i	1			GPRSHLYCRSDYQQRS\TSVCG
		1		1		CMPAVDCCCVLLIHNILRTVM
		1	Ì			WTKYLVTQAVPAR*PGCIRCKC
23613	53981	A	23748	.1	1101	
23614	53982	Α	23749	5	205	YCSSLSFTAAITSRAFLPDRRITT
		1		1		PPATSPSPFSSAMPRRISGPIWTF
		1				ATSRR*TGTPFSLAFKTM
23615	53983	Α	23750	I	761	
23616	53984	В	23751	56	2188	
23617	53985	Α	23752	627	1272	
23618	53986	A	23753	448	2218	VRRQRSDPERSDARMVRFCNL
		1		1		YI*RKNPFILH*LFR*TLRQTKPD
		1		i		NSAGKCVKI
23619	53987	Α	23754	17I	959	AASRGPDAPGLEMAGNCSWEA
		1				HPGNRNKMCPGLSEAPELYRR
		1				GFLTIEQIAMLPPPAVMNYIFLL
i		1				LCLCGLVGNGLVLWFFGFSIKR
	İ	1				NPFSIYFLHLASADVGYLFSKA
		1				LFSILNTGGFMGTFADYIRSVCR
		1		1		VLGLCMLLTG\VLLMPAGWDE
		1				RGVRSERAWCTVETGRVHRSR
		1		ł		VRVSRNTARDPANGLGHLLQS
		1		İ		LGMPALEPGPTLGSALRWEFLQ
		1		1		TASPQLPVQPLHPAAATPQRSL
						RGRGRGPKARALGLGPRSTSYR
23620	53988	Α	23755	664	838	
2362 I	53989	Α	23756	2732	3007	
23622	53990	Α	23757	1522	1706	CHYFACMWSGCEVYS*SAPG**
		1		1	1	C*EDLSGLVRCRCLHSNAETPI*
		Ш.				FAANADTSPRGPFLCR
23623	53991	Α	23758	537	795	ADCPLHMGRPCLID*GPCRTE*P
		1				RCAVVCLLWIAAVSCLSTT/SLR
1						TGI/DGPKYLVTQAVPARYPGCI
		1			į .	RCKCVAVDELASSDMRLPCS
23624	53992	Α	23759	2074	2276	
23625	53993	Α	23760	36	206	ECWIWCELGAELHHQP*ASSH
	i	1		1		WKSSYRQQSSLPRCRASQRTLH
	1	1		1		LLLQGWAVLGPLE
23626	53994	Α	23761	444	725	
23627	53995	В	23762	I	1212	
					<del></del>	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, !=possible nucleotide deletion, !=possible nucleotide insertion)
23628	53996	A	23763	11	972	RFSKSLLDSGVQLASPSGSRTG ATGGAAHSPARAPALHSPWAV DGTRGRLAGTAGLWSPPAFH FKVSLSLPFSCVSSPPGPPLGPS PRKHGMADDAGAGVGGRRPF RGPMGSRGFRGFGSGIRHQ GCGCGQGQGGRGVRGGKAE YKEWMTVTKLGHLVKDIKIKS LEEIYLFSFKTSSSLKDEVLKI MLVQKGTHTSQCTRFAFVAT GDYNGHISLGKCSKEVATAIC GAIILAKLSIVPACRGYWGNKIS KPHTIPCKVAGCCGSVLVCLIS YPRGTGIVLAPCEGDRGLRSS PASLSPESCKDGSHSRSF
23629	53997	A	23764	2	260	LLEECRSLRSRCDQLEERVSAM EDEMNEMK*EGKFREKRISPKR APERSAKHGKEQPVPAAAKSC QNVNTIETRKKLHQLTSKITS
23630	53998	Α	23765	3	1150	
23631	53999	Α	23766	1	1521	
23632	54000	A	23767	1	753	
23633	54001	Α	23768	1	3351	
23634	54002	Α	23769	1	1740	
23635	54003	A	23770	1	1353	
23636	54004	Α	23771	1	1932	
23637	54005	Α	23772	1	1221	
23638	54006	A	23773	I	1214	MAKKQNRKTGNSKKQRASPPP KECSSSPATERWYENDFDELR EEGFRQSNYSELWEDIQTKGKE VENFEKTLEECITRITNTEKCLK ELMELKTKAGLEKECRSLRSR CDQQGERASVLEVLEVLARQL GRKRK*RVFN*EKRKSNCPCLQ TT*LV1*KTPLSQPKISLS**ATS AKSQDTKSMYKNHKHSYTFIT *ESNLQES*NWISSLHLIQKSIQD GLKT*TLDLKPSKP*KKT*ALPF *T*AWARTSCLKHQKQWQQRT KLTNWFLN*HSTLLYGRAN URQLQ*TOTNLQEKNKQPHQKV GEGHQQTILKRRHLCSQKTHE KMSITTGHQRNANQNHNEIPSH TSYNONH

SEQID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
22620	154007	+	23774	11157	2085	OCOMPOS A DECETTIVO DEPLACA
23639	54007	A	23114	1157	2085	QGRPTDFADFGTTIKQDFRLLG
		1			ļ	QTSVDRLLQLSQGQAVKGNQL
		1				LPVSLVKRKTTLAPNTQTASPR
					l	ALADSLIAAGTTGFPTGKRAVS
			ł			ATQLMTQFSPAREQSWRKN/DF
		1	l			DELRKE/GFARSNYS\EKCLKEL
	l					MELKTKARELREECRSLRSRCD
				l		QLEERVSAMEDEMNEMKREG
			1	1		KFREKRIKRNEQSLQEIWDYVK
						RPNLRLIGVPESDAENGTKLEN
			1	1		TLODIIOENFPNLAROANVOIOE
		1				IQRTSQRYSSRRATPRHIIVRFT
		1				KVEMKENMLRAAREKGWVTL
1		1				KGKPIRLTADLVAETLQARRE
		1				WGPIFEEP
23640	54008	Α	23775	1	1707	
23641	54009	Α	23776	1	1899	
23642	54010	Α	23777	1	2781	
23643	54011	Α	23778	1	831	
23644	54012	A	23779	1	3264	
23645	54013	Α	23780	1	882	
23646	54014	Α	23781	1	1853	
23647	54015	A	23782	1	550	
23648	54016	A	23783	1	831	
23649	54017	В	23784	64	3436	
23650	54018	Α	23785	1	1422	
23651	54019	A	23786	1	773	MGKKQSRKTGNSKKQSATPPP
					1	KERSSSPATEQSWTENDFDELR
				ŀ		EEGFRRSNYSELQEETQTKGRE
		1	ļ		Į.	VENFEKNLDECITRTTNTEKCL
			İ		l	KELMELKAKAWELREECRSLR
						SRCNQLEERVSVMEDEMNEMK
		1				REGKFREKRIKRNEQSPQVIRD
		1	1			YVKRPNLRLIGVPESDGKNGTK
						LENTLQDIIQENFPNLARQANIQ
						IQEIPRTS/QRYSSEEHSIIGLIGR
		1				FPS*RTDMFRAPKKREOKRNNR
		1				KIQKEEGGRTAPTGI
23652	54020	Α	23787	ī	2775	
23653	54021	A	23788	1	2712	
23654	54022	Α	23789	1	1590	
23655	54023	Α	23790	I	547	MGKKQNRKTGNSKKQSASPPP
	1	1	1			KECSSSPATEQSWTENDFDELR
	1	1	ŀ			EEGFRRSNYSELREDIQTKGKE
	1	1				VENFEKNLEECITRITNTEKCLK
	1	1				ELMELKTKARELREECRSLSSR
1	1	1				CNQLEERVPAMEDEMNEMKE
l	1	1				D/GNRYDDRRQ/RKEEIMIKSW
!	1	1	l			RFCCRHLIRVVRGWGPSKYLWI
	İ		١.	1		TMVVTLAGNPO
23656	54024	A	23791	1	710	
	1	1				

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23657	54025	A	23792	ı	895	
23658	54026	Α	23793	1	1938	
23659	54027	Α	23794	1	1413	
23660	54028	А	23795	985	1219	LCSGSVERCTESYLHSSHCLVQ QAGVTGDEGGSCY*DPPSHHRS HPRKLKKLCEQEFSVTSAPWRK CRSLHGRFPGLP
23661	54029	Α	23796	I	3570	
23662	54030	A	23797	1	634	
23663	54031	A	23798	I	4551	
23664	54032	А	23799		4721	MHPSQRTPYQSRSPATIFATVST FKSHKSNLETKHPQQVPLSSLG LAANISSEDIESEAEDNGKWES VDRTSFFPALTSFQQSMIHKKI MTVFKSVEPEDGLGRPTLAFHE DRCQLSIQNSLDAGGGCPSMA APPAEVHELASGLISAAALLLSP AASGRVSKHQHIPTGFPDPFKA LHKGKLTTRKDIVTENPSVHHH HQRPKVDKTTKMGKKQNRKT GNSKTQSASPPPKECSSSPATEQ SWMENDFDELREE

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ĺ				sequence		
23665	54033	A	23800	1	2152	MDILGWASRAHKWKGRATGT
ĺ						HVFNLMREYLISLQCCASEQAA
ĺ						SPEHGASFRVWKSLGFQKLILS
ĺ						LICPQCLQSLTRETWMRFKSKK
ĺ		1	1			TKGPDLRLVMDANKWEFRRH
ĺ		1				ALRPALTFPNEGGDLGVEALPV
ĺ		1				ALPPEAHVGGVVLPAALVRGA
ĺ		1		1		HGEQLPAIPGPEEFALPAVQAR
ĺ	1	1				AARPERQVVVIVVNEAGVDPA
ĺ		1		1		GVEGLAPVPVLPGVRLEATAPK
ĺ		l				QPPRLPARLRAPLPSGRKLNAK
ĺ		1				CPRLRWGFFVQDGDGTELACL
ĺ		l				SEGDGGCCQEAPLDHRPRRLRG
ĺ		l				LAGSAGAREAREPRLRPGDWG
ĺ		1				NRIARVRAPEEAAAALSRPARV
ĺ		1				AAALAKTHRREHRARSQKLVM
ĺ		l				SSVQQGKLTTRKDIYTENPSVH
ĺ		l				HHHQRPKVDKTTKMGKKQNR
ĺ		l			İ	KTGNSKKQSAF\PPPKECSSSPA
1		l	ŀ			MEQTWT\DNDFDELREEGFRGS
ĺ		l				NYSELREDIQTKGKEVENFEKD
ĺ		l				LEECITRITNTEKCLKERMELKT
ĺ		l				KARELREECRSLRSRCDQLEER
ĺ		1				VSAMEDEMNEMKREGKFREK
ĺ		1		1		RIKRNEQSLQEIWDYVKRPNLR
ĺ		1				LIGVPESDVENGTKLENTLQDII
ĺ			İ			QENFPNLARQANVQIQEIERMP
ĺ		1				QRYSSRRATPRHIIVRFTKVEM
		l		1		KEKMLRAAREKGRVTLKGKPI
		1		1		RLTAVLSAETLQARREWGPIFN
				1		ILKEKNFQPRISYPAKLSFISEGE
		1		l		IKYFIEKQMLRDFVTTRPALKE
23666	54034	Α	23801	1	2109	

SEQ ID NO:	of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23667	54035	A	23802		2544	MSVHSWKPSWIRPIRSLSPKRRI GTOGTSEVPDTHICOTKERISLSPKRRI GTOGTSEVPDTHICOTKERISLS SGPLLLSVHPVGKLSAVVGRVF HIILSOGRERRRRAVIASRPSEVPG SSHQCVPDSGCRSSVGAHPVREP KQGEALPHSGSPREGKLTNRKD HITKNPSVHIHHHORPKVDKT KIGKKQNRKTGNSKTQSASPPP KECSSSPATEQSWMENDTDELE REGFRÇSNYSELRKDIQTKGKE VENFEKNLEEGKTRITNTEKCL KELMELKTKARELCEECKERS RCNQLEERVSVMEDEMNEM REGKFREKNIKRAELCEECKERS RCNQLEERVSVMEDEMNEM REGKFREKNIKRNEGSLQEIWD VYKRFNLELIGPYESDVENOTK LENTLQDIIQENFPNLARQANV VYKRFNLELIGPYSSRATPHHIV RFTKVEMKEKMLRAAREKDRS TRQKVNKDTQELNSALHQADL LIDIYRTLHPKSTEVTFSAPHHT YSKIDHIVGSKALLSKCKRTEII TYYLSDIISAIKLELRIKNLTYYQNI NYLLINSLLLNDYWYNNEM KABIKMFFETNENKILLELRIKNLTYYQNI KABIKMFFTNENKILLELRIKNLTYYQNI KABIKMFFTNENKILLELRIKNLTYYQNI KABIKMFFTNENKILLELRICECT KUKINESSSWFERINKIDR LARLIKKREKNQIDTIKNDKG UTTOPTEIQTITREYYKHLYAN KLENLEEMDTFLSFIVILNIN,PI

3126

SEO ID	SEO ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
			00000		Lone	New Property Control of the Control
23668	54036	Α	23803	1	1886	MKENFKHRHVVAFISSQEAQE
						DSREEEDRLESRFYFLFLLELSY
		1				KTTGLNREPTVSLNSKSGTPMD
		1		İ		QNEHSHWGPHAKGQCASRSEL
						RIILVGKTGTGKSAAGNSILRKQ
						AFESKLGSQTLTKTCSKSQGSW
						GNREIVIIDTPDMFSWKDHCEA
			l			LYKEVQRCYLLSAPGPHVLLLV
		ĺ				TQLGRYTSQDQQAAQRVKEIF
İ				l		GEDAMGHTIVLFTHKEDLNGG
		ł				SLMDYMHDSDNKALSKLVAA
						CGGRICAFNNRAEGSNQDDQV
						KELMDCIEDLLMEKNGDHYTN
1						GLYSLIQRSKCGPVGSDERVKE
		l			ľ	FKQSLIKYMETQRSYTALAEAN
		l				CLKGALIKTQLPKVDKTTKIGK
		l				KOSRKTGNSKKOSTSPPPKERS
						SSPAMEOSWTENDFDELREEGF
						RRSHYSELPEDIOTEGREVENFE
	Į ·					KSLEECITRITHTEKCLKELMEL
i						KTKARELREECRSLRSQCDQLE
						ERVSAMEDEMNEMKREGKFKE
1						K\RIKRNEQSLQEIWDYVKRPN
1						LRLIGVPESDAENGTKLENTLO
l						DRIOENFPNLAROANGFCRFRN
						HHOTGFSPAGANORGPLAATLS
						GPGGEGOSAVARLTGEKKNHP
1						GAQYANRLSPRVGRFINRSWH
						DRFPDWKAGTGLKSAIFAWE
23669	54037	A	23804	1	4134	Did 10 HE IS I SERSAH AWE
23670	54038	A	23805	1815	1849	
23070	134038	ΙΑ.	23003	1013	1047	

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for tast amino acid of peptide sequence	*-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
	sequence		09/540,217	sequence	or pepude sequence	detection, v-possible nucleotide insertion)
23671	54039	A	23806	1	4016	MGKKQNRKTGNS\KTQSAFP\P
		1				PKECSSSPATEQSWMENDFDKL
					1	REEGFRRSNYSELR\EDIQTKGK
		l				EVENFEK\NLEECIT\RITNTEKC
						LKELMELKTKGSRNYVKECR\S
		ŀ				LRSRCDQLEERVSAMEDEMNE
			1			MK*EGKFREKRIKRNEQSLQEI
						W\DYVK\RPNLRPIGVSESDGEN
						GTKLENTLQDIIPENFPNLARQA
						NVQIQEIQRTPQRYSSRRATPRH
					İ	ILVRFTKVEMKEKMLRAAREK
		l				GQVTLKGKPIRLTADLSAETLQ
		1		8		ARREWGPIFNILKEKNFQPRISY
						PAKLSFISEGEIKYFIDKQMLRD
		Ì				FVTTRPALKELLKESLNMERNN
ŀ		l			l	WTSKQIIQEDGYSEEEGWLCRA
ŀ		l				VIYSHTLQAIMAIVKATGNLQT
		l				DFANPFHTDDARQQFAMSYTV
		1				VGQGALPEDLSRCHLQTEAVQ
		1				RHPESLGCPLDLLHGMPVSSCF
						GYPQDTSSTV
23672	54040	Α	23807	2059	3153	
23673	54041	Α	23808	6	188	
23674	54042	Α	23809	3	1014	VACRPLPRRRAQPATFIPRVAV
		l				KMALVASVRVPARVLLRAGAR
		l				LPGAAFGRTERAAGGGDGA\RP
						\FGSQRVLVEPDAGAGVAVMK
İ		1		1		FKNPPVNSLSLEFLTELVIS\LEK
		l			1	LENDKSFRDVILTSD\RPGVFSA
					1	GLD\LTEMCG\RSPAHYAWVLE
						RPVQELWLRLYQSQL\VLVSAI
	1					NGA\CPAGG\CLVALTCDYTIL\
						AGNPRYCIGLNE\TQ\LGIIAPF\
1		1	1		1	WLKDNPWRTPIGHRAAESA\LQ
		l	1			LGLPLPPPAEALQVGIVDQ\VVP
					1	\EEQVQSTALSAIAQWMAIPKP
					1	MAHK\LTKAIMRKGQGPPAWS
1		1			1	RKGYADVQNF\VSFISKDSIQKS
	L	L	L			LQMYLERLKEEKG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23675	54043	A	23810	11	2665	MTTSIROFTSSSSIKGSSGLGGG
23073	34013	ľ`	23010	ľ	2000	SSRTSCQLSGGLGAGSCRPGSA
		l				GGLGSALGGSSYSSCYSFGSGG
						GYGSSSFEGVDGLLVGGEKAT
l		l				MONLNDRLASYLDKVRALEEA
				i		DTELEVKIRDWYOROAPGPAR
				i		DYSQYYRIIEELQNKPPPSDDLT
		1		1		TLPSPLILTATVDNANILLQNDN
1		1		1		AHLAAADFRTKFETEQALCLSV
1		1				EADINGPCRVLDELTLARADPE
						MHIENLKEELAYLKKNHEEMN
	1	l				APRGQVGGEINVEMGAAPGVD
		l			i	LSRILNEMREOYEKMAEKNRK
		l				DAEDWFFSKTEELNREVATNSE
		i				LVOSGKSEISELRCTMOALEIEL
ļ						QSQLSMKASLEGNLAETENRY
İ		l				CMQLSQIQGLIGSVEERLAQLL
		l			ł	CEMEQQNQEYKILLDMKMRLE
		1				LEITTYHRLLEGEDAQTVQHTP
						GSPCVPEWAHDRCSLPAOLDE
						WSCOAORGICGHRRENCWERIF
						SOSCFYRPTARSLTSLLGTMST
						CSHQFTSSRSMKGSCSIGGSIGG
						GSSRISSVLAGGSCHAPSTYGG
1						ASCGLGGGYGGGFGSSSSFASG
		ŀ				FGGGYGGGLGAGFGGGLGAGL
	1	1				GGGFAGGDGLLVGSEKVTMOH
		1	1			LGDRLASYLDKVRALEEANAD
						LEVKIRDWYQRQRPSEIKDYSP
					}	YFKTIEDLRHKIIAATIENAQPIL
		l	İ			OIDNARLAADDFRTNYEHELAI
	1					OOTVEAGVNGLCREMLALRGO
23676	54044	A	23811	222	548	TLPCGDHHHSODPARLPSWHD
25070	5-10-1-1	l.,	23011		5.0	LLOERSEKCGSQDWQRDSVQP
				1		GVSYOFCCVPWPAGRSPOPRH
	1		İ			CWGCCGTEASYDSGI*SLSTPG
1		1		1		GGQLQGSV*GPDGAGLQNSHR
23677	54045	A	23812	2	165	SSQLQSS. S. BONGBQ. ISSUE
23678	54046	A	23813	I -	665	
23679	54047	В	23814	ti –	1673	
23079	D404/	lo	23014	11	10/3	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	coden for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23680	54048	Α	23815	1	331	ARQLRTSAMTMPVNGAHKDA
		1				DLWSSHDKMLAQPLKDSDVEV
						YNIIKKESNRQRVGLELIASENF
		l				ASRAVLEALGSCLNNKYSEGYP
		ı				GQRYYGGTEFIDELETLCQKRA
		l				LQAYKLDPQCWGVNVQPYSGS
		l				PANFAVYTALVEPHGRIMGLDL
						PDGGHLTHGFMTDKKKISATSI
		l				FFESMPYKVNPDTGYINYDQLE
		l				ENARLFHPKLIIAGTSCYSRNLE
						YARLRKIADENGAYLMADMA
		l				HISGLVAAGVVPSPFEHCHVVT
		1				TTTHKTLRGCRAGMIFYRKGV
		1			}	KSVDPKTGKEILYNLESLINSAV
		1				FPGLQGGPHNHAIAGVAVALK
		1				QAMTLEFKVYQHQVVANCRA
		1				LSEALTELGYKIVTGGS/CVISG
l		1				GAGKPG*FSHQDGTGTSQDGG
						RRPFLHHTLSY
23681	54049	A	23816	3	1709	SHPTPRETPTGVEGQATPSVLG
						AAPASQAKATPTLGRPHASDR
		1				GEVAQLDEWSCQAQRGICGHR
						RENCWERIPSQSCFYRPTARSLT
		l				SLLGTMSTCSHQFTSSRSMKGS
						CSIGGSIGGSSRISSVLAGGSC
		1				HAPSTY/GGGLCVSFSRFSFGGA
		l				CGLGGGYGGGFGSSTNFWS\GF
		l				GGGYGAGLGAGFGGGLGAGL
						GGGFAGGDGLLVGSEKVTMQH
		l				LGDRLASYLDKVRALEEANAD LEVKIRDWYORORPSEIKDYSP
						, ,
		l			,	YFKTIEDLRHKIIAATIENAQPIL OIDNARLAADDFRTKYE\HELA
		l				LRQTVEADVNGLRRVLDELTL
		l				ARTOLEMOIEGLKEELAYLRKN
1		l				HEEEMLALRGQTGGDVNVEM
		1				DAAPGVDLSRILNEMRDOYEO
		l	i			MAEKNRRDAETWFLSKTEELN
		l				KEVASNSELVOSSRSEVTELRR
1		l	ĺ			VLQGLEIELQSQLSMKASLENQ
1		1				L\EETKGRYCMQLSQIQGLIGSV
1		l				EE\QLAQLRCEMEQQSQEYQIL
						LDVKTRLEOKYCHPTAALLEG
1		l				EDAHLSSQQASGQSYSSREVFT\
		1				SSSSSSRQTRPILKEQSSSSFSQ
23682	54050	A	23817	2	365	0000000KQTIG IEKEQ000170Q
23082	D4030	14	2301/	<u> </u>	303	

	SEQ ID NO:		SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23683	54051	Α	23818	1	464	RRWRICSTWTRSFATAWLPPAR
						SANFPRTDRSQGDRGAPAGFAL
		l				APILEFFLWDTSSQKFLQKPHCF
		l				MHGRERAKMGRRAQQESAQA
		l				ENHLNGKNSSLTLTGETSSAKL
		1				PRCR/QGGWAGDSVKASKFRR
		l				KASEEIEDFRLRPOSLNGSDYG
						GDIP
23684	54052	Α	23819	384	805	
23685	54053	Α	23820	3	392	MKYSAIQTLDGEIDLKLLTKVL
		l		1		APEHEVR*VQWWQQFPGLSTL
						AIPINPAELPGLALCLP*RPWNG
						REKVSLQLREKALSLSYPAKPF
		1				LIEDDVGWDWDHLFTEVSSEV
						LTE\WDPLQTEKEDPAGQARHT
23686	54054	A	23821	63	785	GPIRGPWGPGFGGWGGATPPAF
		1				TSLRRLRAAGLAPRRRAKMGR
		1				RAQQESAQAENHLNGKNSSLT
		l				LTGETSSAKLPRCRQGGWAGD
	l	1				SVKASKFRRKASEEIE\DFR\LRP
		ł				OSLNGS\DYGGDIPI\IPDLEEVQ
						EEDFVLQV\AAPPSI\QIKRVMT
						YRDL\DNDLM\KYSAIQTLDGE
						D/IDLKLLTQK/VLAPEHEVRER
		l			ł	NPSWQDDVGLGTWGPICSTEV
		l			1	SLRRSFTEWDP\LQTEKEDPGG
					1	Q\ARHT
23687	54055	Α	23822	1	441	MSFNCSTRNCSSRPIGGRCIVPV
		1				AQVTTTSTTDADCLGGICLPSSF
		l		l		QTGSWLLDHCQETC/CGCQPLG
		l				GISSVCQPVGGISTVCQPVGGV
		1				STVCQPACGVSRTYQQSCVSSC
						RR\PAKCVGASERIKTPRPASCF
		1				QDLPACCLSLNSSS
23688	54056	Α	23823	1	538	PTRPLQQATCSPLSDIMSFNCST
						RNCSSRPIGGRCIVPVAQVTTTS
			l	1	1	TTDADCLGGICLPSSFQTGSWL
		l	l	1	l	LDHCQETCCEPTACQPTCYRRT
			l	1		SCVSNPCOVTCSROTTCISNPCS
	1	1		1		T\PTAGRSP/CVSSGSQPLGGISS
		1				VCOPVGGISTVCOPVGGVSTVC
	[		l	l		QPACGVSRTYQQSCVSSCRRTC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:		Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23689	54057	Α	23824	832	1885	AEWDAADKAHVGPGRSPRQPL
						PHVGRRRAAHTTGRGRRTAGC
		1			İ	APAPARTLCILSILGPWCVLPHL
		1				PTLCLGQAQAASCSQGAHGCF
		ı				AGCSCT/GTSASHPV/CCKCPLH
		l				PVCTLADVHGCITSGFSVISLPQ
		l		l		TCAPWTSWRKWSLSDSWQVD
		l				ACPESCCEPPCCATSCCAPAPCL
		1				TLVCTPVSCVSSPCCQAACEPSF
1		1			i	COSGCTSSCTPSCCOOSSCOPA
		l				CCTSSPCQQACCVPVCCKPVCC
		1	İ			VPVCCKPVCCKPICCVPVCSGA
l		1				SSSCCQQSSRQPACCTTSCCRPS
						SSVSLLCRPVCRSTCCVPIPSCC
1		1				APASTCQPSCCRPASCVSLLCRF
			İ		TCSRLSSACCGLSSGQKSSC	
23690	54058	A	23825		3097	MDEKRTRLOPEEADVPAYPKV
23090	54056	<u> </u> ^	23023	ľ	5077	ATVPQSEKKPSKYPLCSADAAV
		1				STTPPGSPPPPPPPPPPPVLKVLS
		1				SLKPAAPSPATAATTAAAASTA
l		1				ASSTASSASKTASPLEHILQTLF
1						GKKKSFDPSAREPPGSTAGLPQ
						EPKTTAEDGVPAPPLLDPIVQQF
		1				
		1				GQFSKDKALEEEEDDRPYDPEE
	1	l				EYDPERAFDTQLVERGRRHEVE RAPEAAAAEREEVAYDPEDETI
		l				
		l				LEEAKVTVDDLPNRMCADVRR
23691	54059	A	23826	67	442	NSVERPAEPVA VKFFCLYLQSNEKPMKNFKQG
23091	34039	I <sup>A</sup>	23820	0/	442	SNNQLLMRSFWLLCGIDYKKPP
		l				
1		l				PPPAPGPESVAPALSGPPPPAHS RGPRAVVRA\PPPPGPRHRSPAE
1		l				
		l		i		PPPPPGTEPSVATYSIKASKRVS
22.602	54060	١.	02007	-	660	QQDRCHNLMKSQE
23692	54060	A	23827	3	657	GERSRRWGPACGPGAAEKTPP
1		l				CDAGAAGAAPGAPGHAGNPGL
İ		l		İ		GSGPDYD\GEATCMHLHPDDPC
Ì		l				PAGAGCCPGSFEVAFPDAAEK
1		l				MKKSLHSLKKGCYSRVCDLPL
		l				DCPGCDSDSGRQAMFSCIVNFQ
ŀ		l				LPKEESPIPGSSQEESSGPSHSEC
ľ		l				DSVGVDWRGQRVSRKLQAPNH
1		1	1	I		SLVLLVTSLFQETTKSHLNRTN
		L.	L			DAVSPRKFKRLGALYWGKVEA
23693	54061	Α	23828	141	246	
23694	54062	A	23829	1	609	
23695	54063	Α_	23830	1619	1752	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first eodon for peptide sequence	codon for last amino acid of peptide sequence	*-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23696	54064	Α	23831	54	470	CSTMNPSEMQRIAPPRRQRHRS RAPSAHKMNRMVMSEEQMKL PSTKKAEPPTWAQLKLTQLA KKKLENTKVTQTPENMLLAAL KTVSTVSAGVPSSSEESDHRER AMMTTVVLSKRRGKCGEKKEI SDCYCVYVERS
23697	54065	A	23832	213	572	IQWYPYQDTIDFLHRIRKNFFKF HMEPKKSLHCQDNPKPKEQSW MHHTT*LQTMLQGYSNQKRMP PHLGSEERLCPAAPSGR*GAPLP GRCAIFQV*SDSLSAGVPNSSEE TATIENGP
23698	54066	Α	23833	296	583	
23699	54067	Α	23834	2452	4557	
23700	54068	Α	23835	3	266	
23701	54069	А	23836	31	499	ARVTAAVSAAAVAKRVWRDG FDCSGYYSLTTILLVQPTKRPIE GRTYADYESVNECMEGVCKM YEEHLK\RMNPINGPSI/THNTTS QLFDFIDVDLADLSCLVYRAUDT QTY\QPYN\RDWIKDEDPTVLLS SGKAQQAREIIVVGSHWGGWG GAWDTGV
23702	54070	Α	23837	3	323	
23703	54071	Α	23838	366	892	PCVYSOFPAGEQCIKLHSPAEA SPPALEASEAQTRKAPECGSGV PEVAGGSCPCLLLCLPRHQAVP TQGFGTYSPCTSHQPHIFPRPAA PHHLGLLQNPINAASCIQCLYP AGVATTMPRRKAEGDAEGD/K AKWKDEPQRRPAKLSAKPAPPN PEAKPKNAPGVTLSIKGTATRF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nuclcotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23704	54072	A	23839	201	1412	LWKSRLTFKLAMSRVPSPPPPA
23/04	34072	^	23039	201	1412	EMSSGP\VAESWCYTOIKVVKF\
						SYMWDPSITLAFCREEMG\EV\I
				l		KSSTFSSGANDKLKWC\LRVNP
						KG\LDEESKDYLVTFTCYLVSC\
		1				PKSEVRANFKFSILNAKGEETK
						AMESQR\AYRFVQGKDWGFKK
						FIRRDFLLDEANGL\LP\DDKLTL
						FCEVSVVQDSVNISGQNTMNM
		l				VKVPECRLADELRGLWENS\RF
						TDCCLCVAGHEFQAHKAILAA
						RSPVFSAMFEHEMEESKKNPSL
			ŀ			KSMDV\EPEVFKE\MMCFIYTG
						KAPNLTKMA\NDLLAAADKYA
					l	L\ERLKVMCEDALCSNPV/SVEN
						AAEILIL\ADLH\SADQLKTQAV
						DFINYHA\GDVLETSGWKSMVL
						SHP/HNLVAEAYRSLAS\KQCPF
						LGTPTQNALKQSLKILLCC
23705	54073	Λ	23840	I	575	FRHRKIIY*NLLKQEDSINVTEM
						TSIRASVDILIPDPPPKDDEMDT
				l		DKQEKKEVPKCGFLPGNEKVL
						SLLAMVKPEVWTLKEKCIVVIT
		1				WIQHLIPKIEDGNDFGVAIQEK
		1			ļ	VLERVNAVKTKVEAFQTTIS/K
						MDYRALVHERDEAAYGELRA
						MVLDLRAFYAELYHIISSNLEKI
						VNPKGEEKPLMYPTRD
23706	54074	A	23841	1	1092	
23707	54075	Α	23842	130	397	EGVHSQVNLFPIKGNQTHVPAS
						HGTDSALPPTLQMDYRALVHE
						PD*AAYGELRAMVLDLRAFYA
						ELYHIISSNLEKIVNPKGEEKPF
23708	54076	Α	23843	20	334	VRLPGSTHAGGDRSSMAKPCG
					l	VRLSGEARIOVEVLRONLFOEA
					l	EEFLYRFLPOKIIYLNOLLOEDS
						LNVADLTSLRAPLDMPIPDPPP
					l	KDDEMETDEOEKKEGK

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SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last eodon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23709	54077	A	23844		898	MKQKVAAEAGRGGII.QAKVLP SGTLSRLAEALFGVQTYDQDA YGFLTPSAQSMKVFCCLWKLV CKQLEGDVAQGLAIDGDVKKH SGLEWQVETIQNLVDSYMAIV NKNMQDLMPKTIHLMINNTKE FIFSELLANLYSRGEQNTLMGE SAEQAQGREMBREMHHVLRE APSIIGDIKTTIVSTPTGSPKSSPT TQCRLPAVPPARPGLRSPAPGPL PAASTLGEAPVVPSTPGASPDPL GPPPQVPSRPSRGPQPQDRMPC GLAGDLTYGFLEPTLQAIIRWS LLLRPPANSFEQS
23710	54078	A	23845	99	429	FSDAAAVARQPSFGKALGAPQP ADTQPTPSMPPRCPGGSETKPK KAAGKDKSSDK\KNKTKKQMK AEKGGKEKQAEVADQETKDLS AENGDTENKESPASDEATEKDT KSD
23711	54079	A	23846		390	TWMFIAAVFRIASTWKQPRYFS VGKQDQKRTSRKLQQTCSRGA *WLEGKLTNRKE*HQYQRKGH PRKNTIQRSPTLKTKANIILNGQ KLEAFPLKTGTRQRCPLSPLLFN MVLEVLARSIRQEKEIKVFK
23712	54080	A	23847	522	733	ASAADTQANRVWSGPP/RKLQ QTCS*GS*LLQGKLTNRKDIHT KTPSVHHHHQRPKTLNQQRSK ETKKAIT
23713	54081	Α	23848	46	330	
23714	54082	A	23849		873	MAEKKDPTKEPEKKHLLA WM NKLLVELDPKYNDSNTDGSA VYHKIIIACKPVSPPPSAMIGSF PRPSSEADAGAMI.CVQPARTKI LGILGLKATSLQYLRLHTGKPL LPKIDESTKMGRNQPKKAENSK NQNASSPPKDHNSSPVREQNR MQNEFDELTEVGFRRDRSSK NSVVIKALPRHLINGGRSQEWR LGIGARDSFKTSGRRGKSPAGP KTTYPEDEQA/GSRPRPIPTRQ ADPRTPPPTRSEKE/AQLSLATD QMCPSLAPLIK*PQES**WKQRI SI*MEHL
23715	54083	A	23850	1075	1350	AIITANVFKFVSIDRGLVCNMEL EGSNCFIDVCSWTSEMKSKGTS RR*KIT*GNSSGSLPRDTQRSGH RNSSLCCERSKYGGSRVDIRRK VF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
23716	54084	A	23851	1	956	MEVLARVFRQEKEMKGIHIGK
		1			1	EEHKLFLFADNMTLCLEKTKDS
						VKKLVEQPSSFGSVDWLCLSSC
						SGSTHTPRPADVSPGSLPGPGQ
					Į.	TFGTREPPQAVSTKEASSSNLH
				i		APERTVAGLTFTTEQVRALEGV
						FRHHQYLGPLERNWLAREMQL
	ł	ı				SEVQIKTWFQNRRMKHKRQM
						QDSQLNGPLSGSLHGPPAFHSP
		1		ŀ		SSGLANGLQLLCPWAPLPGS/L
		1				WLPLGSFWGLRQMDQEALASS
						WASCCGQPLAYHPPRPGSGAH
		1		1		KLGPALSTGPRGLCALPETGDA
1		1		1		FEETPPTTCLPHKRSVQIASDAY
	ŀ	1		1		PEDSAVLFTSWRHLSP
23717	54085	A	23852	1	846	MEAFDANYTFQSGPSAAGLLV
		1				FAGGPLQALCAWVSAVEAAEQ
	İ					WILQPSSFGSVDWLCLSSCSGST
		1				HTPRPADVSPGSLPGPGQTFGT
						REPPQAVSTKEASSSNLHAPER
				1		TVAGLTFTTEQVRALEGVFRHH
	1	1		1		QYLGPLERNWLAREMQLSEVQ
	1	1			1	IKTWFQNRRMKHKRQMQDSQ
				ŀ		LNGPLMGSFPGAPWLSTHHFLA
				1		LAKGLQ\LLCPWA\PLPGSPGCP
		i		1	i	LGSFWGTP\QMDQEALASSWA
					İ	GCCG\QPLAYHPPRPGSGIDLPK
						GTQLSLRLFQDISSTRGDHRSG
23718	54086	Α	23853	3	239	
23719	54087	Α	23854	10	1154	RRAGFGGVRASGAMGTPPGLQ
ļ				1		TDCEALLSRFQETDSVRFEDFT
						ELWRNMKFGTIFCGRMRNLEK
						NMFTKEALALAWRYFLPPYTF
				I		QIRVGALYLLYGLYNTQLCQPK
	1			1		QKIRVALKDWDEVLKFQQDLV
	1				1	NAQHFDAAYIFRKLRLDRAFHF
				l	1	TAMPKLLSYRMKKKIHRAEVT
						EEFKDPSDRVMKLITSDVLEEM
				1		LNVHDHYQNMKHVISVDKSKP
	1	1		1	1	DKALSLIKDDFFDNIKNIVLEHQ
1	1	1		1	1	QWHKDRKNPSLKSKTNDGEEK
1		1			1	MEGNSQETERCERAESLAKIKS
	1	1			1	KA\FSVVIQASKSRRHRQVKLD
1	1					SSPLDSASGQGQVKATRKKEK
	1			1		KERLKPAGRKMSLRNKGNVQN
						IHKEDKPLSLSMPVITEEEENES
1	1	1	1	1		LSGTEFTASKKRRQH

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23720	54088	A	23855	16	447	INNCDKANCSTTCFNGGTCFYP GKCICPPGLEGEQCEISKCPQPC RNGGKCIGKSKSVCEPGCGAH GTCHEPNKCQCQEGWHGRHCN KRYEASLIHALRPAGAQLRQH TPSLKKAEERGDPPESNIYIWVE LPTSGNVLSYTKFI
23721	54089	В	23856	103	1989	
23722	54090	A	23857	3	1295	SGRGYSRHMADYLISGGTGYY PEGGITAQQLFTGAAARDRLT YNNFLILPGFIDFIADKYDLTSA LTWKITLKMPLISSPMDAVTEA DMALAMALMGGIGFIHHHETTE FQANEVRKYKKFERGFIMDPV VLSPSHTYGDVLEGKMRIGFS GUPITETGTMGSKLDSHKQLLC GAAVGTREDDKYRLDLLTQVG OVDVIVLDLSQGNSLYQMAMYY YIKQKYPHLQVIGGNWTTAQ AKNLIDAGVGLHYGMGCSSIC ITQEVMAYSWPQGTAVYKVAE YARHFDHGEHGKALAGASTV MMGSLLAATTEVPSKYFFSDG VWLKKYRGMGSLDALEKSSGS KQYFMEGDKLKIAQGYSSIQ DKGSIQKFVSYLLAGIQHGCQD TOYCNLSVLLSMMYSGERKFE KWTMLAQIKGGVQGLHSYKK
23723	54091	A	23858	770	1008	LAPSPCVRSKGRKESPDSR/RPL WLRPIAPALGPSGRRGLAPDSR LNPHRSLLGTGNYDVNVIMAA LQGLGLAAVWWDRRR

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nueleotide insertion)
				sequence		
23724	54092	Α	23859	I	1489	MTRWSSYLLGWTTFLLYSYESS
						GGMHEECVFPFTYKGSVYFTCT
						HIHSLSPWCATRAVYNGQWKY
						CQSEDYPRCIFPFIYRGKAYNSC
		l				ISQGSFLGSLWCSVTSVFDEKQ
		l				QWKFCETNEYGGNSLRKPCIFP
		1				SIYRNNVVSDCMEDESNKLWC
		l				PTTENMDKDGKWSFCADTSNL
		l			1	GMGVGPTPRHHRRASTRHSRPP
					1	PADPDGGGERPRTADDRQDDA
l		l				TEDDGTRCRGRQTRTQATNAN
l		l				ANHKKSNRTKPPPRHKPNKAK
						RHTKKETPQNT/CGTQDKNPDS
						GRPKLFRRACITSTGTRGELWT
		1				RSREHPAK\QGRRSGKQRLRPW
l		l				DRMLSPDWHRAQRIAQVWKPE
l						QEDGLSIGFGDQPGPHSDTSSR
		ŀ		İ		QQVRRSKATEMHVGGRVDFD
		ŀ				DFVELMTPKLLAETAGMIGVQ
						EMRDAFKEFDTNGDGEITLVEL
						QQAMQRLLGERLTPREISEVVR
						EADVNGDGTVDFEGDIDGAWG
		1				RHGTQGPHWPWGLGDVRGEPL
23725	54093	Α	23860	283	490	GDIWMGTQPKPYHS\PPGPSPN\
		1				PCPPHFKNQSMPSPKSPPKVFNS
		l				FPALTQKVHTSKVSSETRQVPS
						HL
23726	54094	С	23861	82	222	
23727	54095	Α	23862	27	454	VDGRIAGEITRRGSRARPRPGPQ
						CPPGRPGTAMIKAILIFNNHGKP
					1	RLSKFYQPYSEDTQQQIIRETFH
						LVSKRDENVCNFLEGG/LVFVE
					1	TLDKCFENVCELDLIFHVDKVH
						NILAEMVMGGMVLETNMNEIV
						TQIDAQNKL
23728	54096	Α	23863	1	403	RGSRARPRPGPQCPPGRPGTAM
		1				IKAILIFNNHGKPRLSKFYQPYS
		ŀ				EDTQQQIFRETF/HLGSKKEEKG
]						WNFLKGGLLIGGSDNKLDYRH
		1			1	YATVYFVF\GGDFSKRELGFLN
		_				LIQEFGEPLDKCFEKVWELDLIF
23729	54097	Α	23864	215	599	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23730	54098	Α	23865	2	683	GRIAGEITRRGSRARPRGPOCP PGRPGTAMIKAILIFNNHGKPRL SKRYDPYSEDTQQIIRETTHLV SKRDENVCNFLEGGLLIGGSDN KLIYRRYATULYPVECVDSSES ELGILDLQVFVETLDKCFGNV CEL\DLIFHVJDK VHNILAEMV MGGMVLETMNIEIVTPIDAQN KLEKSEAGLAGAPAIRAIVSAV KKNML/REIPRINIGIDSIKIVP
23731	54099	В	23866	94	726	
23732	54100	В	23867	94	I I 46	
23733	54101	В	23868	1	360	
23734	54102	В	23869	1	1371	
23735	54103	В	23870	1	639	
23736	54104	В	23871	280	2391	
23737	54105	Α	23872	3	244	
23738	54106	В	23873	103	285	
23739	54107	A	23874	70	378	MPLRILLMHGEEEEEEEEEE EEEEEEEEEEEEEEEEEE EEEEEEEE
23740	54108	A	23875	5	156	RKKGGG/EREEEGEGEEEEEE EEEEEEEEEEEEEEEEEVVL LTPGTGAEDFKKH

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SEQ ID	SEQ ID NO:					Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			l	sequence		
23741	54109	Α	23876	1	1808	MEVLSIDNLAGHKVFYQKESHS
			l			WKQQKQDLMSGKRLSAANPLF
		1				ALLLPVLWTTGQAPSPGPGGDP
						ILOLWDDAOOMAGSPRTKKKG
		1				CTRNSQAARLSFOSRGGOGSLT
		l				EKMIFEORFGGGKGESRKDAIA
		1			1	IIVVDIVTLQPCRTVEWVPTRPF
		l				RSLQGHFQGGLRRQQMMVME
						GVWVLSCAMDLTSPRDCLPFC
						VSLPVLLLSAPLCSCSTAGGSPV
i					1	TSPHHFPVEGGLGRVRRCYINE
						PVVTTLGWESVCWVSFWLLLR
		l		1		IRYSTSGQAACHAESQELNPRG
ĺ	ļ.	1		1		FGCLGHRNVTWVKCLLPALQP
		ł				EGSVPFLELLEPFIFASASKTFV
						AANLVTWVACFLLVQAAFQLS
		1		1	Į.	EYRQGVGIPVADRTGDAPTLGL
		1				SYEADVAEGHGFFLDQESVIQII
		1				PEPLPOFYLCLLDAELTGVOGE
						PALPSMQIFSSPEAATGVHNDR
		1				GAEHFPVSRQPSAELGLRPMQA
		1				SOERKINFPGPSPGFPPKRONKT
l		l			1	LSRREKKEEEKEEKEEEEEE/E
l		i				GEEEKEKEEEEEKEEEEEEEE
ļ		1				GGEEEEEEEEEEEEEEE
1		1				KWLYOYDPEDKAQSKOWLSR
		1				GGSGPVKAKADWWRAKEIHSL
		1				OPROVIPGIPSM
23742	54110	A	23877	459	636	IPGGSNGLEAEGPPPPPLPVAAV
		1.	2007	1.07		LA/AGSGGLAVASCAFTICALR
	1					WPGVPCSRVPIWTPS
23743	54111	С	23878	183	254	
23744	54112	A	23879	103	335	LGSGDLPWEINPLSSCSLLHEK
						DPPTTSGPQTDRPKEHLINFKSG
l		1				ACYTCRKSGH*AKECLQPGIPIK
						PHLICVGPH
23745	54113	Α	23880	159	306	LGSGNLP*EINPLSSCSLFREEDP
		ł				PTTSGPQTNQPKEHLTNFKSAA
		l				ED
23746	54114	С	23881	80	106	
23747	54115	Α	23882	317	427	
23748	54116	A	23883	353	511	
23749	54117	В	23884	I	320	
23750	54118	A	23885	1	624	
23751	54119	Α	23886	224	355	
23752	54120	A	23887	872	1003	
23753	54121	Α	23888	I	603	
23754	54122	Α	23889	2	214	
23755	54123	Α	23890	459	601	DVDRHVRGSNFHHNEIRSLAAT
	l	1		i		PRAPGTLAQ/GLTDSFPDLLGLA
		L_		L	l	AED

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
23756	54124	A	23891	2	267	QWDGWGARKGMEWEDDLPLE FGRPTASLLSDRPQPNSSRGSDV PFL\PPSLPCHSSACLAATPRAPG TLAQGSLTDSFPDLLGLAAED
23757	54125	Α	23892	72	299	
23758	54126	В	23893	1	1470	
23759	54127	A	23894	134	1276	
23760	54128	Α	23895	2	847	PPPPRVLRTGGGASLTAQAPGT PRRPQEGEPVTPGRSLGLRSSCQ ASVSLLLPVIVACASLEPGRGR AFRDEGAWTTRVSAAVISSQA ALLSSILWSRRLSFSAHTHAFG TVALGEYAFRMFIRLSARLFGIE VTRPTNS/RSLMKVVKLFSEJW PLAKKETYDWYPPHIITY/AL ELMAGRLRFLGTLTEDEAFRDF MGMEAKTDLKEAFVGKGGNP KEREKGKRAAKKGNSCLVPSR RGDPLSSVAERRKLHLLSFPHIG GDVIFLMGSLFGRYTSHLGEI
23761	54129	Α	23898	3	1224	MRRNKPRORGVREARPAGGAG PFWRGARCSGAGEGGGERRG SPPAALALAPASGPRRNFPPDA RCLIYQQIHQGAFLLAGSASLS AVTGYPEGGARTFEDVCPEK MFGFHKPKMYRSIEGCICISGA KSSSSRTDSKRYEKIDFQSCFG LHETRSGDICNACVLLVKRWK KLPAGSKKINWNHYDARAGP SULKTTFEPKIKVKTLSGINRIKS TQISKLQKEFKIRSNSDAHSTTS SASPAQSPCYSNQSDDGDTEM AGSSNRTPVSFLDLTYWKRQK ICCGIIYKGRFGEVLIDTHLFKP CCSNKKAAAEKPEQDEPPLIS LPAGCSLHGLAADWTTDMITV TTSKCREPSCQOEGSFQRKAF PKRLPKMAEVQVLVL
23762	54130	A	23900	69	355	
23763	54131	А	23901	316	830	VSLSQTLHPPMTLQLSRYSRGW GLSSDRPRGAKAKVPSCFGTAP RIGPEAAPRLLAHHLVPAGRGS/ ARAPAPGWRLGSP\GAGRKGR DFRLPASGLAVSGEWDPIPWGY VGEVGGCSCQWFSPGVQQKRY RRKVWLGQPQDLPGPSALADK GPALAKADGQRVNVASVGLQP

de hod				
.	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop eodon, /=possible nucleotide
١	09/540,217	sequence	or peptide sequence	deletion, \=possible nucleotide insertion)
A	23902	I	1044	MSYKVTQEISKELRWISAVLVT
ı				WDLRGETGTRVCIAMGVSRMQ
				YSIMAKIVGSGNNLELNPAADK
ŀ				LDNLGAQCGLASVKVSARAAP
- 1				PFAPATLVALLEARAAELPLPPP
				PGVSLRCVEAPDVRGRGRGGG
- 1				EGSGVTRLLLPPPPPSGASFLQR
- 1	1			GRPVTPALFQAGWGRAADRPG
- 1				PADLAGGAGGRSGGRRMLPRS
				PGAALRDRSALEDGIVEGYGPL
				GEPEEGSRSSFLESKVAFNYCN
ŀ	ļ			KVAPLDFGNEAVEQCHTMSDR
- 1				KAVIKNADMSEDMQQDAVDC
- 1				A\TQAMEKYNIEKDIAAYIKKEF
				DKKYNPTWHCIVGRNF\GSY\V
				PHTEDKHFIYFYLGQVAIL\LFK
A	23903	33	427	
Α	23904	72	777	EWKQAGEGGGGQRRGSPPAVL
				ALAPASCPRREMSYLPTNPPGSI
				LICWLGVSLSSHRSAGGGSAIY
				G\SYYPEEKMFVFHKPKMYRSI
				EGYCICRAKSSSSRFTDSKRYEK
- 1				DFQSCFGLHETRSGDICNACVL
1				LVKRWKKLPAGSKKKTGIMCN
				OSDDGSDTEMASGSNRTPVFSF
				LDLTYWKRQKICCGINYKGRF
				GEVLIDTHPFKPCCSNKKAAAE
				KPEEOGPEPLPISTOEW
Α	23905	234	989	PKTSDOAYYCPEEKMFGFHKP
1				KMY\RSIEGC\CISGAKSSSRFT
				DSKRYEK\DFQ\SCFGLHETR\SG
				DI\CNACVLLVKRWKKLPAGSK
				K\NWNHVVDARAGPSLKTTLK
l				PKKVKTL\SGNRIK\STQISKLQK
				/ENFKR\HNS\DAHSTHPOVPSP\
	1			AOSPLFTVNOFRWTGSDTGVG
	1			FPGSNRNHPVFSFLDL\TYWKR
	1			OKICCGNYKGRFGEVLIDTHLF
- 1				KPCCSNKKA\AAEKPEEQGPEP
1 _				LPISTOEWVTEVFM
A	23906	3	1059	
	A	A 23903 A 23904 A 23905	A 23903 33 A 23904 72	A 23902 1 1044  A 23903 33 427  A 23904 72 777  A 23905 234 989

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23769	54137	A	23907	1	1697	MRPCALCWRSAVTAAPERAVL
		l				ESRGGAPDTPRRLHSFTQTRSLS
						PQAHAGPHTRLHPRQGHAHMR
						GGGRVTDTRRHIPNADVPRYK
						LTCTGGVTTFVALYDYESRTET
		1				DLSFKKGER\LQIVNNTKGDW
						WLAHSLSTGQTGYIPSNYVAPS
						DSIQAEEWYFGKITRRESERLLL
			ŀ			NAENPRGTFLVRESETTKGAYC
		i				LSVSDFDNAKGLNVKHYKIRK
						LDSGGFYITSRTQFNSLQQLVA
İ						YYSMSHCPPFSAAGWYGGGCP
						EEPQKHADGLCHRLTTVCPTSK
						PQTQGLAKDAWEIPRESLRLEV
						KLGQGCFGEVWMGTWNGTTR
						VAIKTLKPGTMSPEAFLQEAQV
		l				MKKLRHEKLVQLYAVVSEEPI
	ľ					YIVTEYMSKGSLLDFLKGETGK
		l				YLRLPQLVDMAAQIASGMAYV
						ERMNYVHRDLRAANILVGENL
						VCKVADFGLARLIEDNEYTAR
İ						QGAKFPIKWTAPEAALYGRFTI
						KSDVWSFGILLTELTTKGRVPY
	ŀ					PGMVNREVLDQVERGYRMPCP
						PECPESLHDLMCQCWRKEPEER
						PTFEYLQAFLEDYFTSTEPQYQP
23770	54138	Α	23908	1	736	MDPGAALQRRAGGGGGLGAG
						SPALSGGQGRRRKQPPRPADFK
		1				LQVIIIGSRGVGKTSLMERFTDD
						TFCEACKSTVGV\DFKIKTVELR
						GKKIRLQIWDTAGQERFSSITSA
						YYRSAKGMILVYDITKKETFDD
						LPKWMKMIDKYA\SEDADLLL
			l			VGNKLDCETDREITRQQGEKFA
						QQITGMRFCEASAKDNFNVDEI
		ĺ				FLKLVDDILKKMPLDILRNELS
						NSILSLQPEPEIPPELPPPRPHVR
23771	54139	В	23909	84	441	
23772	54140	A	23910	6	254	
23773	54141	A	23911	1	230	
23774	54142	A	23912	3	187	
23775	54143	A	23913	3	143	
23776	54144	A	23914	3	481	
23777	54145	Α	23915	1	711	
23778	54146	Α	23916	55	381	

SEQ ID	leen in vo.	Alot	SEQ ID NO:	Nucleotide	Nucleatide losation of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1.0.	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		L				
23779	54147	Α	23917	240	875	GLSAESTPTSIMSMTLGYWDIR
						GLAHAIRLLLEYTDSSYEEKKY
	İ					TMGDAPDYDRSQWL\NEKFKL
	i					GL\DFPNLPY\LIDGAHKMHPEA
						TPFLCYI\ARKHNLLLGETGRRE
İ						KISSWDIL\ENQAMDVSNQLAR
		1				VCYSPDFEKLKPEYLEELPTMM
						OHFSOFLGKRPWFVGDKITFVD
1		1				FLAYDVLDLHRIFEPNCLDAFP
						NLKDFISRFEVSCGIM
23780	54148	A	23918	64	774	CLSAESAPTSTMPMTLGYWDIR
25700		Ι'	23310		l'''	GLAHAIRLLLEYTD\SSYEEKKY
1					l	TMGDAP\DYDRSQWLNEKFKL
1						GL\DFPNLPYLIDGAHKITOSNA
		1				ILRYI\ARK\HNLCGGDRRREKIS
		1			l	CWDILENQVM\DNHMELVRLC
ŀ		1				YDPDFEKLKPKYLEELPEKLKL
l .		1				YSEFLGKRPCFLG\DKITFVDFL
		1				
				i	ļ.	VY\DVLETK\QVFEPK\CLDAF\Q
	ļ	1				NLKDFISRFEGL\EKISAYMKSS\
22701	54140	١.	22010	162	217	RFL\QELVFSKMAVWGNK
23781	54149	A	23919	163	317	LSPNNYYRKLPQKVKLE*RYHR
						LALFSFPTKKLTPEKVSLCCRFL
		Ļ.		10		EMGYIF
23782	54150	Α	23920	62	255	VRLTAQKSLGPLTSTHSLWGSA
		1				\RARHREPVGSVCGCDPCEQLL
		_				LLVSQLRAPGVDSAAAGRPV
23783	5415I	Α	23921	173	467	KSRFCFFGVVLKLPKKLHIR*WI
		1			\	GATL*NQP*TIFQTDGDCSHQSL
						GPLPACDLWDQLHLRSKQGGS
		1				VCGCDPCEQLLLLVSQLRAPGV
		ــــــــــــــــــــــــــــــــــــــ				DSAAAGRPV
23784	54152	Α	23922	35	548	PRSCPSLSPGLSCLLQ/RLVNYQI
		1				SIKCSNQFKLEVCLLNAENKVV
						DNQAGTQGQ\LPAVGQARWGS
		1				GCRGAHTW\PLAVIGAADCTG
		1				QRGWEGV*AFPGLTEPVSSLFC
		1				GQDGD/CSPHSM*VLSHVVLWL
i		1				LNGLC\SLVLGSVCGCDPCEQL
		1				LLLVSQLRAPGVDSAAAGHPV
23785	54153	Α	23923	I	599	MSPAGLVNYQISVKCSNQFKLE
		1		1		VCLLNAENKVVDNQAGTQGQL
		1				KGRPDGAQAV/VGAHTWRFGC
1	1	1	l	1		RRCG*LHRSHWGL*LLHTPCGR
	İ	1				PHCARHREPVAHSLLCSRAISGS
	1	1			1	EPHFWQCSVQ*PQLCVPGRVAT
1		1	l			PLGELDSGDGDCSHQSLGPLPA
1	1	1	l			CDLWDQLHLRSRQGGSVCGCD
		1	l	1		PCEQLLLLVSQLRAPGVDSAAA
		1		1		GRPV
23786	54154	A	23924	290	528	
25,00	124124	1.	23724	200	520	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23787	54155	A	23925	274	571	KSRFCFFGVVLKLPKKLHIR*WI
						GATL*NQP*TIFQTDGDCSHQSL
				İ		GPLPACDLWDQLHLHSRQGGS
1				1		VCGCDPCEQLLLL\VSQLQAPAI
		1				DSAAAGRPV
23788	54156	В	23926	204	410	
23789	54157	Α	23927	29	330	YLGVELVKSAYPFFQIHLSFPSC
		1		1		MWLCGLFLLFYIDFLTLMCGD
		1				KPLICKKHQLSASTCA/CTIC*G
						ALTLLMEIEQGPCDLSCASREV
						STARSLAELFSGP
23790	54158	Α	23928	259	699	ETSHQVMDRSNPVKPALDYFL
						NRLVNYQISVKCSNQFKLEVCL
				•		LNAENKVVDNQAGTQGQLKV
						LGANLWWPYLMHEHPAYLYS
		1				WEDGDCSHQSLGPLPACDLWD
1		!				QLHLRSRQGGSVCGCDPCEQLL
		ļ.,				LLVSQLRAPG\VDSAAAGHPV
23791	54159	A	23929	245	695	
23792	54160	Α	23930	302	745	ETSHQVMDRSNPVKPALDYFS
		1				NRLVNYQISVKCSNQFKLEVCL
				1		LNAENKVVDNQAGTQGQLKV
		1				LGANLWWPYLMHEHPAYLYS
		1	ì			WEDGDCFTPKALDPLPACDLW
		1				DQL\HLRSRQGGSVCGCDPCEQ
23793	54161	١.	23931	2	293	LLLLVSQLRAPGVDSAAAGRPV VFCHTDLRKNLTVDEGTMKVE
23/93	34161	Λ	23931	ľ	293	VLPALTDNYMYLVIDDETKEA
		1				AIVDPVOPOKVVDAARKHG\V
		1		1		KLTTVLTTHHHWDHAGGNEKL
		i .		i		VKLESADLGIRLF
23794	54162	A	23932	1	441	
23795	54163	Α	23933	1	630	
23796	54164	A	23934	3	265	
23797	54165	Α	23935	3	1062	SAVQPGPVMVVGRGCSRRSLA
		1		1		ALGAACARRGLGPALLGVFCH
1		1		1		TDLRKNLTVDEGTMKVEVLPA
		1	ŀ	I		LTDNYMYLVIDDETKEAAIVDP
1		1				VQPQKVVDAARKHGVKLTTVL
			į.	l .		TTHHHWDHAGGNEKLVKLESG
		1	ı			LKVYGGDDRIGALTHKITHLST
			1			LQVG\SLNVKCLATP\CHTSGHI
		1	l			CYFVSKLGG\SEPFVVFTGDTLF
1	1		1			V\AGCGEVLMKGLRDEMCKAL
			1	1		LEV\LGRLPPDTVGSASLRAGR
					1	VYCG\HEYTIKQTSKFAR\HVET
			1	1	1	GNA\AIREK\LPWA\KEKYSIGEP
1	1	1	1	1		TVPST\LAE\EFTYNP\FIKVREER
	1	1	1	1		RCKQH\AGETDPVTTMRPVRRE
		1	1	1		KDQ\FK\MPRDLRPPLHLQQDL
		L.	l			GIKALLDRLF

SEO ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23798	54166	Α	23936	107	750	SHLWSPVSEPLQACSFC/VIRDK
l		i				YRYFACLMRARSEEHTNEKDM
1						MKAIRLLKEAEEEFWYHOHPO
l						PCIFPDSPRGTSYVTHECYKVPE
l						WCLDDWHPSEKAMYPDYFAK
						RKOWKKLRRESWKREVKOLO
İ						EETPPAGKEECKLHEGMNYVV
						LTCIPGAIKQYLACSRYSKTPGP
		l			Ī	RDSCWEVQSDNQCWWWLDSG
		1	1			AHGLVTSFKGVMMTVEATTM
						AAVF
23799	54167	A	23937	3	718	PASTISCSVTRSRRAVSRLSARP
23777	34107	n	23737	ľ	' ' '	GKVSAVMASLASGPYLTHQQK
						VLRLYKRALRHLESWCVORDK
						YRYFACLMRARFEEHKNEODM
						AKATOLLKEAEEE\FWYROHPO
				1		P\YIFPDSPWGHPPYERY\DCYK
						VP\EWCLDDWHPSEK\AMYP\D
						YFAKREQWKKLRRESWEREV\
						KQL\QEET\PPGGSF*LEALPPGP
		1			1	KRKVILPPLWW\YIVTRP\RERP
		1				MLERERPHLSCLQVKYVTEHG
23800	54168	A	23938	269	488	TDGVIFIGLSWSPRRCGLGIORD
23800	34106	^	23936	209	1400	AERIFFALDQASCGFTIHPHIDA
						AETPRGLT\MHLCRRCQRGQSQ
						AEKLP
23801	54169	A	23939	1	1560	MVHLGAPVMALNLOGKHLCD
23601	34109	^	23939	ļ.	1300	STASSLLPRRRESNSDDTARED
						AROKRPANTDMRTOELOTRET
					1	GOGTARETEVATPLSGPSAGDS
ŀ	ĺ					PPAASEMSCGQLVLALLLLFFL
1	ŀ					LVWNTGTKSGSGRVARLAQLA
1				1		HDCDGGLALGLCYAVLDQRSV
1						VQATPQAPKPGPVQPLTVQGL
1						OPVHVAOESSGNEITARROSAR
		l				SRPRGDRTAPSRRVTRRPKHAP
				•		ERAPIQQGNIVTVRAPTARKKW
		1		1		RPAVGTPTGAHPVCTAKGTGG
1						
		1				RPTPTQAHTRYTERDPIRNSAEL TGSRPCTGRERSGVEACKAAV
		1				
						HNLL\AQTGSSGLIINYPLDD\Q
		1	١,			DAIAVEAACTNVPALFLDVSDQ
	1	1				TPINSIIFSHEDDVWRTPLVAPT
	1	1	1	1		NLNTRSSQPARHHRAGAVRRH
	1	1	1	İ	1	HMTVARSATAHARLRRHAKPQ
		1	1		1	GRERIKPSGAREGSEPLGGESGI
1	1	1	1		1	RTPRGRERDPNPSGAKAGSEPG
1		1	1			RIPETLPRCKGETEAKCWARGN
1						FHTAQGSWPRGGRSRWEPLEA
		_				PALSQQAHGGELPLPAA ARSHR
23802	54170	В	23940	98	229	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	
NO:	nf peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23803	54171	Α	23941	1	256	
23804	54172	Α	23942	289	433	VLWSCAAEDFDWLVYRPIGSA
				l		LSCRLIGPWAIGMWETGFP*VL
				l		WSCAAEDFDWLVYRPIGSALSC
	l					RLIGPWAIGMWETGFPLLSSVS
				1		LS
23805	54173	Α	23943	450	1260	LCACVVGGEKRSGPRIGCAGA
						GKDPFWRAVGVAQKAAAVVA
				1		CGAASPYRNRADVSVSDHSQS
						RGPGGLSLSGGTVCDMGKKTK
	1					RTADSSSSEDEEEYVVEKVLDR
	1	1				RVVKGQVEYLLKWKGFSEEHN
		1				TWEPEKNL\DCPELISEFMKKY
	l	ĺ				KKMKEGGN\NKVREKSESSKR
	l	l	(			KSNFSDSAEDIKCKKKREOSND
		l				IARGFERGLEPEKIIGATDSCGD
						LMFLMKWKDTDEADLVLAKE
		l				ANVKCPQIVIAFYEERLTWHAF\
						PEDAENKEKETAKS
23806	54174	Α	23944	3	267	AMVGGGGVGGGLLENANPLIY
						QRSGERPVTAGEEDEQVPDSID
				1	1	AREIFDLIRS*AWPLTPQPAYW
						YPGPSSCQGCYRCFLESLTMRN
23807	54175	Α	23945	1	942	
23808	54176	Α	23946	3	470	AMVGGGGVGGGLLENANPLIY
1		١				QRSGERPVTAGEEDEQVPDSID
l	1		1			AREIFDIRRCWARAGSGGLRW
	1					GEQ*YRAAGGAASQQGVPGRG
	1					F*VGKGAESLFLHFPQ\LIRFLN
	1	l				DP\EHSLTLEELNVVEQVRVQV
		1				SDPESTVAVAFTPTISHWSMGT
23809	54177	Α	23947	2	604	ARSHRISGGGSAMVGGGGVGG
		l				GLLENANPLIYQRFGE\RPVTAG
	1					EEDEQVPDSIDAREVF\DLIRSH
						QMTPEHP\LTLEELNVV\EQVR\
	1				į .	VQVSDP\ESTVAVGFPKPTHSGT
	1					CRHGPPLIGLS\IKVK\LLRSLPS
	1	1			1	AFSRWDVAHLLPGDPLPQEHA
	1	1		1	1	VEQATLQIKEAGWATALGRTH
		1				HLLEVVN\QCLSSPAPGTWAFV
23810	54178	Α	23948	1	1002	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Aminn acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
	<u> </u>	<u> </u>				
23811	54179	A	23949	89	2587	STSRDRRLSKRSRTTMAAQPRG
		1				GDYRGSFLLSILLGTPWEAWAG
						RILYSVSEETDKGSFVGDIAKDL
						GLEPRELAERGVRIISRGRTQLF
İ						ALNQRSGSLVTAGRIDREEICA
		l				QSARCLVNFNILMEDKMNLYPI
		1				DVEHDINDNVPRFLTEEINVKI
		l				MENTAPGVRFPLSEAGDPDVG
1		l				TNSLQSYQLSPNRHFSLAVQSG
		ł				DDETKYPELVLERVLDREEERV
						HHLVLTASDGGDPPRS\$TAHIQ
		1				VTVVDVNDHTPVFSLPQYQVT
				1		VPENVPVGTRLLTVHAIDLDEG
1		l				VNGEVTYSFRKITPKLPKMFHL
						NSLTGEISTLEGLDYEETAFYE
1		1		ł		MEVQAQDGPGSLTKAKVLITV
		l				LDVNDNAPEVTMTSLSSSIPED
		ŀ				TPLGTVIALFYLODRDSGKNGE
						VTCTIPENLPFKLEKSIDNYYRL
					1	K\TTKNLDRETLSLYNITLKATD
					i	GGTPPLSRETHIFMQVADTNDN
1					ł	PPTFPHSSYSVYIAENNPRGASI
		l l				FLVTAQDHDSEDNAQITYSLAE
		l				DTIOGAPVSSYVSINSDTGVLY
		1				ALQSFDYEQLRELQLRVTAHDS
		1		l	İ	GDPPLSSNMSLSLFVLDQNDNP
		1		1	İ	PEILYPALPTDGSTGMELAPRSA
		1				EPGYLVTKVVAVDKDSGQNA
	i	1				WLSYLLLKASEPGLFAVGLYTG
		1			1	EVRTARALLDRDALKQSLVVA
		1				VODHGOPPLSATVTLTVAVAD
1		l				SIPEVLADLGSLEPSDGPYNYDL
23812	54180	В	23950	165	335	SILE VENDESCRET SPOT TIVIDE
23812	54180	A	23950	217	859	GSAMAQQRALPQSKETLLQSY
23813	J#181	l^	23931	21/	0.55	NKRLKDDIKSIMDNFTEIIKTAK
		ŀ			İ	IEDETQVSRAT\QGEQDNYEMH
						VRAANIVRAGESLMKLVSDLK
						QFLIL\NDFPSVNEAIDQRNQQL
		1	1			RTLQEECDRKLITLRDEISIDLY
		l				ELEEEYYSSSPGPSLVSQQLSEQ
1	1	l	1			QLLSSAWTAWPKVTLSSLAPGP
1		ŀ				QYSVNGPAAIIAHPGTCASAGIS
		L_				SSLLPEPVLLVPVF
23814	54182	Α	23952	1	1263	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino aeid	*=Stop eodon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	İ			sequence		
23815	54183	ΙΛ	23953	70	2530	L PGVSRLVSKLRRARGMGSGAG
23813	34183	A	25955	l' <sup>0</sup>	2550	
						ELGRAERLPELFLLTLFCPAL
						CEQIRYRIPEEMPKGSVVGNLA
	İ					TDLGFSVQELPTRKLRVSSEKP
	į.					YFTVSAESAELLVSSRLYREEIC
	İ					GKKPACALEFEAVAENPLNFY
	1		l			HVNVEIEDINDHTPKFTQNSFEL
						QISESAQPGTRFILGSAHDADIG
						SNTLQNYQLSPSDHFSLINKEKS
	1	l				DGSKYPEMVLKTPLDREKQKS
	ļ.					YHLTLTALDFGAPPLSSTAQIH
						VLVTDANDNAPVFSQDVYRVS
	l					LSENVYPGTTVLQVTATDQDE
	ì	l				GVNAEITFSFSEASQITQFDLNS
	į.					NTGEITVLNTLDFEEVKEYSIVL
						EARDGGGMIAQCTVEVEVIDE
	Į.	1				NDNAPEVIFQSLPNLIMEDAEL
		1				GTHIALLKVRDKDSRHNGEVT
		1				CKLEGDVPFKILTSSRNTYKLV
1		1			1	TDAVLDREONPEYNITYTATOR
		1				GKPPLSSSSSITLHIGDVNDNAP
						VFSQSSYIVHVAENNPPGASISQ
	1	ł				VRASDPDLGPNGQVSYCIMAS
	1	ł				DLEQRELSSYVSISAESGVVFA
	1	1				ORAFDHEOLRAFELTLOARDO
		1				GSPALSANVSLRVLVDDRNDN
						APRVLYPALGPDGSALFDMVP
	l .					HAAEPGYLVTKVVAVDADSGH
	l .	ŀ		1		NAWLSYHVLOASEPGLFSLGLR
					İ	TGEVRTARALGDRDAVRORLL
						VAVRDGGOPPLSATATLHLVF
		ł				ADSLOEVLPDITDRPDPSDLOA
23816	54184	С	23954	75	149	The state of the s
23817	54185	A	23955	442	618	CORLRSWVSOPFSFSFFPSFPTF*
23017	34103	1^	23733	1772	010	NILCLFVYLEVEFPSCCPGWSN
						NWAKSSGKALEAN
22010	64106	<del> </del> -	22056		245	INWARSSURALEAN
23818	54186	Α	23956	1	245	l

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, ≔possible nucleotide insertion)
23819	54187	Α	23957	30	1079	APCASFLTPLAAASVGKVWLG
						LRGAAPMASVLSYESLVHAVA
						GAVGSVTAMT\VVFPLDTASIF
					į.	DFSVDF\KRKSKTTHMVVLEIIK
				1		EEGLLAPYRGWFPVISSLCCSNF
						VYFYTFNSLKALWVKGQHSTT
						GKDLVVGFVAGVVNVLLTTPL
						WVVNTRLKLQGAQFRNEDIVP
						TNYKGIIDAFHQIIRDEGISALW
						NGTFPSLLLVFNPAIQFMFYEGL
			l	1		KRQLLTFAFQLSSLDVFIIGAVA
			ŀ			KAIATTVTYPLQTVPLFPQFGR
	ľ		l			HRLNPENRTLGSLRNILYLLLQI
						FRTPSEREPFYTSVTLSVLSSVA
						YPMMPLSNITSAYRIGVLEKGG
			l	ĺ		GSGKSDNSDDDDDDDDDGGV
23820	54188	A	23958	1375	2451	VPRVPFAINVLTHSRQRYRRNL
						RFYADDPTIRV/GGPTYHWVRE\
						AEEERVVDNRMHDRFCELRTA
						AGHPVEGGRPLVIKGITHPAET
						ALQRFKFLMYQVVASDLDGTL
			l			LSPDHTLSPYAKETLKLLTARGI
						NFVFATGRHHVDVGQIRDNLEI
						KSYMITSNGARVHDLDGNLIFA
						HNLDRDIASDLFGVVNDNPDIIT
						NVYRDDEWFMNRHRPEEMRFF
						KEAVFQYALYEPGLLEPEGETK
						ALLTPGKLKNFLVEKMRTLGT
	1					AACPPYHIAFVIGGTSAETNLKT
	1					VKLASAHYYDELPTEGNEHGQ
		l				GSAMSSWNRNCWKRPRNSVLA
		l				RSLAVNTSRTTFALSVCHVTAH
		1				PARSAWASPVR
23821	54189	В	23959	1	1386	
23822	54190	Λ	23960	1375	1609	VPRVPFAINVLTHSRQRYRRNL
						RFYADDPTIRV/GGPTYHWVRE
				1		S\EEERVVDNRMHDRFCELRTA
						AGHPVEGGRPLVIK
23823	54191	Α	23961	433	741	
23824	54192	Α	23962	12	224	
23825	54193	Α	23963	3	2053	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
23826	54194	A	23964	1	2408	MGGADIAGPRREHNAKHPTVG
	1		1			LGSAAQTTCLPGPKQKGDPTRH
			i			QPVDKRNDVGPRQHFWCWRV
			l .			EIHKEKRMKTGPLNESELEWLD
						DILTKYNTDHAILDVAELDGLL
						TAVLSSPQEIEPEQWLVAVWG
1						GADYVPRWASEKEMTRFMNL
			l			AFOHMADTAERLNEFPEOFEPL
i						FGLREVDGSELTIVVPPGGRTA
	İ					TNALSCGSLFGEETSPRGPSADS
1						TSOSIGGGSPHDCAKGIALVAA
	İ					NGGDIRDYEGVDRSAKPOLPMI
			ļ			AINTTAGTASEMTRFCIITDEAR
						HIKMAIVDKHVTPLLSVNDSSL
1						
	ŀ					MIGMPKSLTAATGMDALTHAI
						EAYVSIAATPITDACALKAVTM
	1					IAGISTFFASSRMALMQASAPSA
						SFLPVTFTPIAAAQSRRRAAAT
			1			GYRDRWLCHGDVLLQICANTQ
						DTVIHALRDIIKHNSCPDDCSVQ
į.						HDAVIDDGCDFIDYLGGSETVV
1						ALNTPRADDYHPVMVLCAKTD
1						KAHRNVIHTSKLCFITLTPEISQ
						RSHGKSSKRVFSCFPVFLLLKH
						RRIVAQKYAELAYDLFHLGFD
						VLIIDHRGQGRSGRLLADPHLG
						HVNRFNDYVDDLAAFWOOEV
						QPGPWRKRYILAHSMGGAISTL
	i		1			FLORHPGVCD\AIALTAPMFGIV
						IRMPSFMAROILNWAEAHPRFR
1						DGYAIGTGRWRALPFAINVLTH
			1			SRORYRRNLRFYADDPTIRVGG
22027	64106	<u>_</u>	22077	,	22.62	SKOKTIGGER TADDITIKYGO
23827	54195	В	23965	14602	3357	CICDODDI BODI DA A TOTOLO
23828	54196	A	23966	14682	14931	EIGPGPRPLPSPLP*ATSTSVLAA
			1			SGRPERTR\HAGIKIVLEDIFTL
						WRQVETKVRAKIRKMKVTTKV
	l					NRHDKINGKRKTAKEQSPLLQE
1	l	1	1			SLFATGDVSHNLLRALDVGLLA
1		1	1			NLSALAELDISNNKISTLEEGIF
			1			ANLFNLSEINLSGNPFECDCGL
			l			AWLPRWAEEQQVRV/AAARGS
1	1	1	1			HVCWAWLPGWPASAWHPLAG
			1			QWLW*GVCRLPP*QQLRHRGS
1	l	1				SVLFSCPRRPA/CSQRPAAPSAS
						PPARASQPSPSRA
	L	L			L	TTTT TO TO TO TO TO

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
23829	54197	A	23967	1	714	MVKLSIVLTPRFLSHDOGOLTK
25025	15.1.57	ľ.	120,00	1		ELOOHVKSVTCPCEYLRKVINT
		1				LADHRHRGTDFGGSPWLLIITV
		1			l	FLRSYKFAISLCTSYLCVSFLKTI
		l				FPSONGHDGSTDVOORARRSN
		l			İ	RRRQEGIKIVLEDIFTLWRQVET
		l			ĺ	KVRAKICKMKVTTKVNRHDKI
		l				
		l			ł	NGKRKTAKEHLRKLSMKEREH
	1	1			i	GEKERQVSEAEENGKLDMKEI
	1	ı	l			HTY/MITPSARKPLCNRVRMAA
		_				AEHRHSSGLPYWPYLTAETL
23830	54198	Α	23968	111	216	ALLFPISTWSTPLGTQQAPQCV*
						PTPMPMRPVPS
2383I	54199	A	23969	I	4710	
23832	54200	A	23970	3544	3723	RSHRSPPRCPPSRAPAPRRSAAA
ŀ	i	l			İ	PPRR*PPRAPAAPRPGPP/CMAR
1	ł	ı				PPPARPTAGSRVPPL
23833	54201	A	23971	3	1119	EFGTSHRHRGTDFGGSPWLLIIT
	1	ı				VFLRSYKFAISLCTTYLWVINTL
		i			i	ADHRHRGTDFGGSPWLLIITVF
		ı		ŀ	1	LRSYKFAISLCTTYLC\VSFLKTI
	l					FPSQNGHDGSTDVQQRARRSN
		l				CRROEGIKIVL\EDIFTLWROVE
	1	l				TKVRAKIRKMKVTTK\ATRLTK
		l				IKERRKTAODHWRKLSMKERE
		l				HREEEROVSEAEENGKLDMKEI
		l				HTYMEMFORAQALR\RRAED\Y
	l	l				YRCKITPSARKPLCNRVRMAAA
	l	1				EHRHSSGLPYWPYLTAETLKNR
		1				MGHQPPPPTQQHSIIDNSLSLKT
		l			1	PPECLLTPLPPSALPSADDNLKT
		1				PAECLLYPLPPSADDNLKTPPEC
	l			1		LLTPLPPSAPPSADDNLKTPPKC
		1				VCSLPFHPORMIISRN
23834	54202	A	23972	1	1149	VCSLFFRIQKWIISKN
23034	34202	lv.	23912		1149	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
110.	sequence	nou.	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		ĺ.
		<u> </u>				
23835	54203	A	23973	93	1165	HRRSAAEAPAAAVMPFPFGKS
						HKSPADIVKNLKESMAVLEKQ
1			l			DISDKKAEKATEEVSKNLVAM
l						KEILYGTNEKDPQTEAGAQLAQ
1	1	l				ELYNSGLLITLVADLQLIDFEGK
	1					KDVAQIFNNILRRQIGT\RTPTV
l	1					EYICTQ\QNI\LFMLLKGYESP\EI
ĺ	1					ALN\CGIMLRECIRHEPLGKIIL
i	i					WSEQFYDFFRYVEMSTFDIASD
						A\FATFKGLLTRHKLLSAEFLEQ
	1					HYDRFFSEYEKLLHSENYVTKR
			l	1		QSLKLLGELLLDRHNFTIMTKY
			l			ISKPV\NLK\LMMNLLRDK\$RNI
			1			QFEAFHVFKA\FVANPNKTQPIL
	1					DILLKNQAKLIEFLSKFQNDRTE
						DEQFNDEKTYLVKQIRDLKRPA
						QQEA
23836	54204	Α	23974	3	158	
23837	54205	A	23975	3	1071	GVRASGRAVPRAVFAGMKRPC
						EETTSESDMDETIDVGSENNYS
ŀ						GQSTSS\VIRLNSPTTTSQIMAR
						KKRRGIIEKR\RRDRINNSLSELR
l			İ			RLVPTAFEKQGSAKLEKAEILQ
l	1				l	MTVDHLKMLQATGGKGYFD\A
l	1					HALAMDFMSIGF\RECLTEAAR
						YLSSVEGLDSSDPLRVRLVSHL
					ľ	STCATQREAAAMTSSMAHHHH
	1					PLHPHHWAAAFHHLPAALLQP
						NGLHASESTPCRLSTTSEVPPAH
						GSALLTATFAHADSALRMP\ST
l	1	l	1	1		GSVAPCVPLLSTSLLSLSATV\H
l	1	l	1			AAAATATAAAHSFPLSFAGAFP
l	1	1	1			MLPPNAAAAVAAATAISPPLSV
	1					SATSSPQQTSSGTNNKPYRPWG
l	1		l	1		TEVGAF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23838	54206	A	23976	1	1416	MPGNLTRKESSSPLTDRHCHTS
						SQFSHRFNAISPFPPAELGCDHV
		l		l		ASAAADPLGVPGVATQWAGGL
		l		l		SAALAAPRRPHSCRRLOVLGAL
		l				GILKSAGRVNFRCFGPLSIRGKP
		l		ł		KTPLWEQEKPKLQTGSAGSQQ
		l				KRVMLKPSRGADRVLGRGSQR
						PAETDRETTDLRQRKRMKEGK
						ERTLRAARGETRRSNGNLARO
			ì			TVPEGCRSQTQALQVVRNART
		1				PPCIPNLDRPPEACSSGSTGGM
				i		MPLRVAPSPVASGESSHLSSGR
		l		l		GDRGLVRCKVEELRFQPVAFPE
				1		FIPPPTSPRADSESGDWCCTVLA
						NPSKTGSGGWEQRKTEC\PQGR
		l				TONPOORGS/RNLVLLTAGOES
						TPRRSQLTRSPAAERRTEEQGE
		1				DRKNRALGCNLFPRVPRHARD
		1				SORALRASTOAPKKSTFAFPGI
						HSGAVCSKRRICGVQGEPFPLPF
						ASQCSSCLKILPSPGPTKELTSY
						LLHSGCGFLDKLRV
23839	54207	Α	23977	3	139	
23840	54208	A	23978	141	734	GSQLRRPAFGVRVALEESKGLI
						RKDSCRGRACAGANACARRRR
l		1				PPTMSGAAWRGTPLALVRSSRT
t		1				RGVWAWQWSATRSPSDSCFPG
		1				VARDKEDGEPKTKHLREEEEE
		1				GEKHRELRLRNYVPEDEDLKK
						RRVPQAKPVAVEEKVKEQLEA
			1			A\SPEPVIEEVDLANLAPRKPDW
l				1		DLKRDVAKKLEKLKKRTQRAI
						GRLDP
23841	54209	Α	23979	3	585	GTVGGVCGPRRSSSASPQEQDQ
					ł	DRRKDWGHVELLEVLQARALF
		1				TAVPLSEEDRATVLKAVHKVK
					1	PTFSLPLQLPPPVNTSKLLRDVY
		l				AKDGRVSYPKLHLPLKTLQCLF
		l				EKQLHVELASRMCVVSVEKPT
		1				LPSKEVTHARKT\LKTLRDQWE
1						KALCRELRETKNRLERQVYEG
1	I	1			I	RFSLYPFLCLLDECEVVRMLLQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23842	54210	Α	23980	1	3778	MRRRGPVGWPGAACVMSALC
		1				WGRGAAGLKRALRPCGRPGLP
		1				GKEGTAGGVCGPRRSSSASPQE
		l				QDQDRRKDWGHVELLEVLQA
		l				RVRQLQAESVSEVVVNRVDVA
		l				RLPECGSGDGSLQPPRKVQMG
		1		1		AKDATPVPCGRWAKILEKDKR
		1				TQQMRMQRLKAKLQMPFQSG
		ł	l			EFKALTRRLQVEPRLLSKQ\MG
		l				GCLEDCTRQAPESPWEEQLARL
		1				LQEAPGKLSLDVEQAPSGQHSQ
						AQLSGQQQRLLAFFKCCLLTDQ
23843	54211	Α	23981	101	805	HMYTSMKPSSQPRSWISTPHPK
						ACSSLSFPLSLDNWMNINLREVI
						CTSRDGDKFWRMPECYIRGSTI
			1			KYLRIPDEIIDMVKEEVVAKPR
		1				GRGGLQQQKQQKGRGMGGAG
		1				RVGSPLSSG\VPGAGRQPHVAS
		l				LLLGVALAALPRRLPNLPGPPS
		1				CPLAMNRGPGRGAPRHGTQPSS
		1		1	l	QCASWRSSGHIKSPPQAPVSPM
	1	l				RGNGGAHLSRPLATTLGTLRGS
				ļ. <u>.</u> .		IQSQGRPPPPEMMNWH
23844	54212	A	23982	133	1069	RIAWGQDRSCLQGGPFATYVP
						GLILCSLQGWYGVQAGQDLQG
	Ì			1	İ	LPRSQGWARRPGGKLITCSPSP
					1	QDGDKFWRMPECYIRGSTIKYL
		i				RIPDEIIDMVKEE\VVAKGR\GR
		1				EGPCQQAESQQK/GVRGMGGA\
		1		1		GRCVFGGRGRGMGSPITG\RCH
		ł		1		PDKKPGRQAGQTVSAHPDRLL RPLLARVGDFRSTVSERNLLOK
		l		1		GRGRPPRGEPPFRVLVTPSLFKV
	1	1		1		
	İ	1		1		KAPFFLPKNRQILRVRKCLLCCF FGFCETGPSOKLGWNPLERVLC
		1			1	LKAIFSSPFASGFSRASHFPRRG
	i	1			l	LFLHFLGSLSEASNOTGRFLILK
23845	54213	A	23983	1	326	PVSLECYTVPPEDNLALLQLYF
23043	54213	۱^	23703	1.	1320	RTLVTGA/LRPRWCPVLYAVTL
	l	1		1	1	WPHVNSFIFSQRPRIVDIDGKM
		1				GTLSCIOPFNRCLARRRTLCPNG
	1	1		1		ROETKEOKACLPGSRLFELF
23846	54214	A	23984	2	341	NYL TREORACLE OSKLIELE
23840	134214	I <sup>A</sup>	22904	14	Dat	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		<b>!</b>		<del>                                     </del>		
23847	54215	A	23985	114	712	DLISALMIWFHVIVIHTPYVCIYI
		l				PTLPLSTSFPLSQLPVSLECYTV
						PPEDNLALLQLYFRTLVTGALR
		1				PRWCPVLYAVAVAHVNSFIFSQ
						DPQSSVSYISIFRGRKGASN*AA
		ļ.				SGPAGEEVSVKQGQWWVPAL
		l				GSVPPLPVLLL\DEVKAARRSM
	İ	ŀ				LQKTWLLADEGLRQHLLHYKL
						PNSTLPEGFELYSQMPPLRQHY
23848	54216	Α	23986	357	2771	FESSCPPVGSPVGAGPTPSLYKV
						PCATAMKLLRVLASAGRNIAA
		1				RLLSSFDLRSRLCRIIAEAPQEL
						ALPPEEAEMLSTEALRLWAVA
						ASYGQGGYLYRELYPVLMRAL
						OVVPRELSTHPPOPLSMORIASL
		l				LTLLTQLTLAAGSTPAETISDSA
	l	l				EASLSATPSLVTWTQVSGLQPL
		l				VEPCLRQTLKLLSRPEMWRAV
		l				GPVPAACLLFLGAYYQAWSQQ
		ı				PSSCPEDWLQDMERLSEELLLP
		l				LLSQPTLGSLWDSLRHCSLLCN
		l				PLSCVPALEAPPSLVSLGCSGGC
İ		l				PRLSLAGSASPFPFLTALLSLLN
1		1				TLAQIHKGLCGQLAAILAAPGL
		1				QNYFLQCVAPGAAPHLTPFSA
		ļ.			<u> </u>	WALRHEYHLOYLALALAOKA
						AALOPLPATHAALYHGMALAL
	l					LSRLLPGSEYLTHELLLSCVFRL
		1				EFLPERTSGGPEAADFSDQLSL
		1		İ		GSSRVPRCGQGT\LLAQACQDL
ŀ		l	į.		1	PSIRNCYLTHCSPARASLLASQA
					l	LHRGELQRVPTLLLPMPTEPLL
		l	į .			PTDWPFLPLIRLYHRASNTPSG\
		l				LSPTDTMG\TAM\RVLQWVLVL
			i			ESWRPOALWAVPPAARLARLM
1		l			ŀ	,
1		1				CVFLVDSELFRESPVQHLV\AAL
1	l	1	ł			LAQLCQPQVLPNLNLDCGLRG
1					1	LTSFPDLYANFLDHYEAVSFGD
					1	HLFGAMVLLPLQRRFSVTLRLA
		l			1	LFGEHVGALRALSLPLTQLPVS
						LECYTVPPEDNLALLQLYFRTL
23849	54217	Α	23987	291	454	KAETGSLPYTLCKNQFKMD*RL
			1			TC*T*NHKNPRRKPR*YHSGHR
		_				HGQRLHDENT
23850	54218	Α	23988	1	613	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /-possible nuclcotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23851	54219	A	23989	807	1345	RCEERLLPLNLAVIALPSGLER
		1				WSFFAVYNGHPGSQVAKYCCE
l		l				HLLDHIADNQ/DFTGSAGAPSLE
		l				NVKNGIRTG/CPEPEVHDTERSE
		l				EDDHFIILACDSIWDVMGNEEL
		l				CDFVRSRLEVTDDPEKVYNEIV
		l				DTCLHKGSQDNMCVIF\LWFPN
						APKVSPEAAKKEAELGKYRLN
		l				SILSDQ
23852	54220	A	23990	344	636	SSLSLLDIWVGSKSLLL*IVPQ*T
		l				YVCMCLYSSMIYSPLGIYPVMG
	İ			i		WLGQMVFLVLDPSGIATLTST
	i	l				MVELVYSPTNSVKVFLFLRILSS
		l				TCCFLTF
23853	54221	Α	23991	l	2793	
23854	54222	A	23992	18	409	SSVVEFVDPRVAPALLPQLPLES
						SSHGSVLPTMAVCLDKGHKMT
		l				KNMSKPRHSCRRGRLTKH\KFM
		l				WDMIREVCGFAPYERHAMVLL
		l		İ		KVSKDKRALKFIKKRVGTHIRT
	1	1				KRRQEELSNVPAIMRKADAKK
						D
23855	54223	A	23993	1	609	MHPRGPPPPAQEESHTDESPGID
l		1				HQSLARRAGRVGRTGGFSQLL
		1				AQTNTSMLSSSPASCTSPSPDGE
		l				NPCKKVHWASGRRRTSSTDSES
l						KSHPDSSKIPRSRRPSRLTVKYD
		1				RGQLQRWLEMEQWVDAQVQE
	1	l				LFQDQATPSEPEIDLEALMDLST
						EEQKTQLEAILGNCP\RPTEAFIS
		I				ELLSQLKETPETQPASEISLRDH
23856	54224	Α	23994	I	204	
23857	54225	A	23995	1	418	MWNRLCNWVTSIGWKSLEGSE
l						EDRKSSQWGTLPSGAVRRGPPS
	1					SRSKNGRSNSSLHPVPGKATGT
	1			1		QLEPMRAVMGDESCKATGTEL
	1			1		LHALGAHPLHRLLKQHSHMMF
	1	1		1		QKVQVEPLRK\EPHHWQRL*PF
	1					SSE*NHGPSR
23858	54226	A_	23996	104	253	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		·
23859	54227	Α	23997	666	1436	WRPCFWLQAKKEAPAP\PKAK
23003						AKVKAL\KAKKAVLKDVHSHK
		1				KKKD\RKSPTFRG\PKTLCTR\KP
		l				PKCLQKTAPRKNKLDHCAITKF
		l				PLTTKSTTKKIEDNNTLVFIVDV
		1				KANKHOFKOAVKKLYDTDVA
		l				KNITKRWDVTSIITSCYIHLRLA
		l				SRLALEDLLAGLMNCLKGTFW
		1				NEESGERVETRPESRFKQQSKLI
		1				YTIKNPGRKKYGAEEKRAFWC
		1				HTDLLLHDHDEQVNHFSSPSST
		l				FTKCKRKKPFLLREFVWIK
23860	54228	A	23998	2	301	FIRCERORFFEEREFVWIR
23861	54229	A	23999	1	242	MGMVRKQALERTEWALVCRQ
23001	J 122)	ľ`	23,777	ľ	1	AWSSRMGYSTTVSPTOCOYTC
						SRGLLDSOLOLYKCTOALPEIH
		l				SLGHVCPSLITVKYSHYVLAAA
				ł		TLFIWDNNESLRTGLLSVSTCK
						SGTRDFMRISGMVSLTSEVDNV
		l				EVFHYFPLHIQGKKFWAVGAI
	1	1				MTIFVISIYYEPGAHWHHMWK
						LLKVYSLESQKREEDISKRLPNF
		1				LKSSRRPWEDLILMLRKSPEMS
		1				KEYRASPLSLSAALRSKDRQSR
		1				HRCVTSTFHIREVRCCKSQGLP
		1				LCMHETAAHICP*TEGOOKGSG
		1				GLLYIP*TSPEIFSASESDPPMAC
23862	54230	Α	24000	2	948	CGTPTVVPGTGTSRRVRTATVE
						PTWKCRAAQQPPPPP/PPLWVK
			ŀ		ł	LPPPDTSTKPELQLEELDADSS
	i			l		WMPLHGSCPTGKLAVDARSSL
	•	l		l		FTEPAFPYWKKMKHTFQNILS
	ŀ	l				WSKFTNLTYSVTQLWLLVVMA
	İ	1				QTSYPHSLVGRKQVILHSSCLY
		l				SLDNKTSLAGDDHDDDDHDDN
		l				DSSPAEKGPPPITKSLDKADEQS
						HLRLAVLELGFYKPVNLLLVA
		1				MTGCLLQLMMKKEQCGTHGG
		l	i			EGSTLEGSRPDVPQDLFSPTLGF
		1				YGSTSLSADLGENVILKLYLPSF
						PLQESEPSAEHGENLWALYVGF
						EERRSEASLAKAP
23863	54231	Α	24001	265	513	
23864	54232	Α	24002	513	796	TGTAWTPPPPPLTTGAPCTPPPR
	1	1		I	1	CTARGRT/PGDSHLGGGPAATA
	1	l	1	1	1	GGPRTSPMSSGGPSAPGMRPPA
				1		SSPKRNTTSLLNSGLEPTFSFRIT
			L			FGFM

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *-Stop coden, /-possible nucleotide deletion, \-possible nucleotide insertion)
23865	54233	A	24003	1268	2097	TTELTGAISLSTLGDPSPFRQPS AQMGHLLSTSLSTPPTISSSYLPT QPAFATVTPSSSSVLPGLPQTSF SGMGPSAELMASTISPKQQLPQF SAAFGHQLSSHSGIPKDLQPSHS SIAPPTGFTVTGVKLQTFAVSVT VLKATCLELFVPPGGFVVSLAS GVKLQTVTPLHFSGNHPSAFYY YRLSFEMDKGKKRQFGVEKEK CGSLGGFVMLGNEBMIRKAV LLAWKLIPFLGIREAKNPGSENI RLATILEAPPPSWELWEQGPPG APYYQP
23866	54234	A	24004	3	1163	GEVTSWNRGGGLRIWRYFLEGD GVOKTMELLISRLTYCWLEGDG RVSWRLGLEIGSLPTPVPVRLCP ANKSPPMVCGTTGRSSNSSTST KRMCTCSSRAALIFILKGPEGG TLMEEIRALIKEPPESSLTPSTRG GHSEKAPSKNHQAATVNTTLIS RISHISHTHLLIHCTGKEQHEPY EDAATRSRPEADLSRHERLTCR HLDLGLPDMONRQCVGSGGL RPVGGLEGPRHWCLVRSSRLG VLPRRPQGCEDDAGGLQRQDC GQWRELVQLPTPPVCPGGASP HPKPLPDLPDPPDSSRCPCLLPG TCPAAPPAGSPSPSCSSPCCLLPG VPTPSVLECCTQVLLEDSKPQS HRNSAPHLQPGIWSHIPGSHPW CLVLAAEGSTSSV

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23867	54235	A	24005		2502	MEARVERAYOKROVLFLCVFL GMSWAGAEPLRYFVAEETERG TFLTNLAKDLGLGVGELRARG TRINSDONMOILLISSLTGDLLL KEKLDREELGGPREPCVLPFQL LLEKPFQIFRAELWYRDINDHA SYFLDREISLKLIESTTGAAFLL ESAQDSDVGTNSLSNYTISPNA YPHINVHDSGGGINYPELVLNQ VLDREEIPEFSLTLTALDGGSPP RSGTALVRILVLDVNDNAPDFV RSLYKVQVPENSPVGSMVVSV SARDLDTGSNGEIAYAFSYATE RILKTFQNPTYGSLHKAQLDY EAIOTYTLTQAKDGGGI,SGKC TVVVDVTDINDNRPELLLSSLT FYTLVTIGAKDGGGI,SGKC TVVVDVTDINDNRPELLLSSLT FYTLVTEKPLDKERNTEVNITIT VTDLGTPRLKTEHNITVLVSDV NDNAPAFTQTSYTLEVRENNSP ALPIGSVSATRDBGSTAGNSP ALPIGSVSATRDBGSTAGNSP ALPIGSVSATREFRNNSP ALPIGSVSATREFRNNSP ALPIGSVSATREFSRNSP ALPIGSVSATREFFRNSP FALHIGSVSTAETNIQVTTSLLP PRNPHLPLASLVSINTDNGHLF RSLDYFMFHLPLASLVSINTDNGHLF RSLDYSSTAETNIQVTTSLLP PROPHLPLASLVSINTDNGHLF RSLDYSTAETNIQVTTSLLP PROPHLPLASLVSINTDNGHLF RARSLDYSALGFEFFRVGASDR GSTALSSEALVRVLVLDANDSS LFVLFPLQNGSAFCHEMPRAVA AEPGYLVTKNAWLS*QLLKAT EFGLFGVWAHNGTDRLLSERD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23868	54236	Α	24006	1	1960	MEARVERAVQKRQVLFLCVFL
		ı				GMSWAGAEPLRYFVAEETERG
						TFLTNLAKDLGLGVGELRARG
		l				TRIVSDQNMQILLLSSLTGDLLL
						NEKLDREELCGPREPCVLPFQL
			ł			LLEKPFQIFRAELWVRDINDHA
						SVFLDREISLKILESTTPGAAFLL
						ESAQDSDVGTNSLSNYTISPNA
						YFHINVHDSGEGNIYPELVLNQ
						VLDREEIPEFSLTLTALDGGSPP
						RSGTALVRILVLDVNDNAPDFV
			1			RSLYKVQVPENSPVGSMVVSV
						SARDLDTGSNGEIAYAFSYATE
						RILKTFQINPTSGSLHLKAQLDY
						EAIQTYTLTIQAKDGGGLSGKC
						TVVVDVTDINDNRPELLLSSLT
						SPIAENSPETVVAVFRIRDRDSG
	1		l			NNGKTVCSIQDDLPFILKPSVEN
						FYTLVTEKPLDRERNTEYNITIT
			1	ľ		VTDLGTPRLKTEHNITVLVSDV
			1			NDNAPAFTQTSYTLFVRENNSP
			1			ALPIGSVSATDRDSGTNAQVIY
						SLLPSQDPHLPLASLVSINADNG
			İ			HLFALSPWTTMPLRAFEFRNNS
			ł			PALHIGSVSTAETNIQVTYSLLP
			Į.			PRNPHLPLASLVSINTDNGHLF
			l			ALRSLDYEALQEFEFRVGASDR
		l				GSTALSSEALVRVLVLDANDSS
						LFVLFPLQNGSAPCTELMPRA\A
23869	54237	Α	24007	1	966	MGHHLMIDKLVALGGLYYAIQ
					ľ	RHYATKCSVLKNDQILVIGLFM
			ŀ			IQNVIYRKHFANPLSALFLQQGI
						ELFAAIAEIHITTVAERNHAITQI
				1		RLEAQTIFDSLKEWHNAIRKSL
						DYEALQAFEFRVGATDRGFPAL
				ĺ		SSEALVRVLVLDANDNSPFVLY
	i					PLQNGSAPCTELVPRAAEPGYL
	l					VTKVVAVDGDSGQNAWLSYQ
						LLKATEPGLFGVWAHNGEVRT
						ARLLSERDVAKHRLVVLVKDN
	1					GEPPRSATATLQVLLVDGFSQP
	l					YLPLPRAAPAQAQADSLTVYL
						VVALASVSSLFLFSVLLFVAVR\
			L			LCRRSRAAPVGRCSVL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			i	sequence		
		<u> </u>				
23870	54238	A	24008	[1	2673	MEVMGNMGSKILITGITSTKRK
						VALLAKKRERKIISQRKDVTDS
1						RAKRQSEEKSIGKETVVIGIGLI
1					i	SKDSAAAICESRWRLSESHFER
						LIQRRMEARVERAVQKRQVLF
	1					LCVFLGMSWAGAEPLRYFVAE
1		l				ETERGTFLTNLAKDLGLGVGEL
						RARGTRIVSDQNMQILLLSSLT
1						GDLLLNEKLDREELCGPREPCV
						LPFQLLLEKPFQIFRAELWVRDI
	1					NDHAPVFLDREISLKILESTTPG
	1		i			AAFLLESAODSDVGTNSLSNYT
						ISPNAYFHINVHDSGEGNIYPEL
1						VLNOVLDREEIPEFSLTLTALDG
l	1					GSPPRSGTALVRILVLDVNDNA
	1	i			j	PDFVRSLYKVQVPENSPVGSM
	l					
1						VVSVSARDLDTGSNGEIAYAFS
		1				YATERILKTFQINPTSGSLHLKA
						QLDYEAIQTYTLTIQAKDGGGL
	İ	1				SGKCTVVVDVTDINDNRPELLL
	i	1				SSLTSPIAENSPETVVAVFRIRD
						RDSGNNGKTVCSIQDDVPFILK
	1					PSVENFYTLVTEKPLDRERNTE
	1					YNITITVTDLGTPRLKTEHNITV
					1	LVSDVNDNAPAFTQTSYTLFVR
						ENNSPALPIGSVSATDRDSGTN
						AQVIYSLLPSQDPHLPLASLVSI
ŀ					1	NADNGHLFALRSLDYEALQAF
1		ļ.		1		EFRVGATDRGSPALSSEALVRV
1	Į.					LVLDANDNL\PFVLYPLQNGSA
}						PCTELVPRAAEPGYLVTKVVA
Í		ľ	ŀ			VDGDSGQNAWLSYQLLKATEP
23871	54239	В	24009	1	993	T D G D G Q T T T T T T T T T T T T T T T T T T
23872	54240	В	24010	i	4056	
23873	54241	В	24011	i	906	
23874	54242	В	24012	li .	4318	
23875	54243	В	24013	41	1935	
23876	54244	Ā	24014	3	328	GWOSGPVLGGLCGTAGOSCGD
200.0		'		ľ	1	GDGIWWAQAAQAGDPAHLRL
1	l		Į.			LTHGQPSSVPEAPVRAHQPALP
						QPRSAAARECGR/AGRSTTNSL
	i		l			RKRVASESYMIVAPPMPSSWSS
	l					
22275	-1015	l	24015	114		SG
23877	54245	A	24015	114	567	VKKPHSAEWVPLWGLQPGWR
		1	ĺ	ł		QPFRCNSGSEKELRRPVSNSQEI
				1		LSPVTMRKPAVSVCCDLTLREA
		1	l	1		APPFQRHERTYTGEKPYKCKQC
1		1		1		GKAFSGFCYTQILEKAQTGKQP
		1	l	I		YE/AHERTHWKQNHDCTQFGK
1		1	l	I		AFSFSSSLCIHEKIHTGEKLYE
	L					

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nueleotide	Nucleotide location of last	Amino aeid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequen <del>c</del> e	deletion, \=possible nucleotide insertion)
23878	54246	Α	24016	170	491	PAQHQRGLWQSPTHQALTLML
						SVHREDWKPVLTINSIIYGLOYL
						FLESKNRWLGLERLGFCPSGLIT
		l				PPTCTGHRSP/NPEDPLNKEAAE
		1				VLQNNRRLFEQNVQRSMR
23879	54247	A	24017	1	1062	
23880	54248	Α	24018	2	700	RRPTRPGGGSGGGGRMIKLFS
						LKQQKKEEESAGGTKGSSKKA
						SAAQLR\IQKDINELNLPKTCDIS
ŀ						FSRSQTTLLNFK\LVICPDEGFY
i		1	İ			KSGKFVFSFK\VGQGYPHDPPK
	l	1	i			VKCETMVYHPNIDLEGNV\CLN
		l				ILRGDWEASPLRLNSIIYGLQY\
1			ľ			LFL\EPNPEDP\LNKEAAEV\LQN
						NRR/VCFEQNVQRSMRGGLHPA
						PPTFERCLKIGLAHTHPPPRPQA
				1		LASPAKYLLGAMG
23881	54249	Α	24019	302	884	PMAPLSSQQPTLFPNTLTQAQQ
						VQLTKGLRKWMVQHGGDGW
						VVEENRTTVPGAPSQTCFVTSF/
						EVNAPAIQSWCCKKQVLDLEE
						EGLWPELLDSGRIEICVSDWWG
		1				ARHDSGCMYRLLVQLLDANQT
ŀ		1		1		VLDKFSAVPDPIPQWEQQCLPF
						TFTHVFFQHSRWGVRFCVFSEH
						RGPGTHSSGAGPLLSPCEQLQC
23882	54250	A	24020	274	866	SSLLNLPSSWDYGWVSYEQAN
	1					CKGEQFVFEKGEYAHWDWWT
						SSQRTDFSSLR/RPSKVDSQEPK
l						V\YLYENPNFTGKKMEIIDDD\V
İ						PSFHA\HGYQEKVSSVWVRSG
		l				MWVG\YKYPGYRGLQYLLKKG
		1				DYKDSSNF\GARHPQV\QSVRCI
		1				HDMQGHQCGAFHPLGTQEEEE
l		1				AGQKEQMEQVPFWLARGDICE
		L				LSGKKSVI
23883	54251	A	24021	179	374	AASIGRSMTEKAHAGPRPGRN*
1		1			1	ADAPSSSDTLFPPDALLSNQAG
						RLACWLFSYGPQTKNDFHIFK
23884	54252	Α	24022	1	141	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23885	54253	Α	24023	Īī	506	PHSPLWKTLCSPSSGVCCGFVT
						LFCNDRFACLFQLTPPPSPAWM
		1	ŀ			TAAASLLASLLPALPPLNLQDA
		l				KFVEERRKQLQNYLRTVMNKV
		l				IQMVPEFAASPKKETLIQLMPFF
						VMGLQVGMYGLFHFGVMMDQ
						PQPSLTRADESPCDRVLSPCAA
			ŀ			HRKQRDCCLSWKAEPE*FLVK*
						KTLCSPSSGVCCGFVTLFCNDR
					<u> </u>	FACLFQLTPPPSPAWMTAAASL
						LASLLPALPPLNLQDAKFVEER
		ĺ				RKQLQNYLRTVMNKVIQMVPE
						FAASPKKETLIQLMPFFVMGLQ
				1		VGMYGLFHFGVMMDOPOPSLT
						RADESPCDRVLSPCAAHRKOR
						DCCLSWKAEPE
23886	54254	Α	24024	1	396	
23887	54255	Α	24025	1	1176	MVPGDLSQTSEDQSLSDFEISN
						RALINVWIPSVFLRGKAANAFH
						VYQQSIAGVSGIARKRALIPFGK
						STFSYVCITNIKLTDALPSWDYT
	ĺ			1		CHFNLPKAQKPGEPTKCILTME
			ł	1		WLALLQGDPGSWIQKEFIVGKR
			ŀ			GGKMDNILWDQRDVQPPGLRP
	l			1		HQLLHSGGGPLISFRNGGPILV
	ĺ		ŀ			WELRRKGASGKEIVGSRYFRES
					1	DEVLVWKNPKTLQKEDEEVNS
		l				THPMHTYPCTHCQVFCVTAEII
		1				NIYLQACIVLEMSMLPPGERPLF
						LLRLTTVLEVTVFDSQWSRPYH
		ĺ				VSADPSAWKHHHHYHHHHHY
		l				QLSSSSPTITIINHPSPS/LSHASPS
	İ		l			P/CHPSPSPAVHHHHQPSITITSR
	1	1		l		PSPSPAVHHHHQPSITIIIRPSSITI
						INHPSQASSIHHHHQPSIPFIVSH
23888	54256	Α	24026	1349	2125	
23889	54257	Α	24027	670	972	KTDEWNIYRRYTEFRSLHHKLQ
		1				NKYPQVRAYNFPPKKAIGNKD
1		1		l		AKFVEERRKQLQNYLRSVMNK
		l				VIQMVPPSSLPAPKKETLIQ\LM
		l	l	I		PLLRSTSTPPRRAL
23890	54258	A	24028	1	699	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			l	sequence		
23891	54259	Α	24029	221	864	VDCPRSFSAVVVWQGRFWLLS
İ						FIPGGFLGMTLFFASGMNWTVT
			l			LLGFLAGNLEVEPSDTIENVKA
			l			KIQDKEG/IPPDQQRLIFAGKQL
				ŀ		EDGGTLSDYNIQKESTLHLVLR
						LRGGMKIFVKTLTGKTITLEVE
	ŀ	1	ł	i		PSDTIENVKAKIQDEEGIPPDQQ
ł		ĺ	ł			RLIFAGKKLEDGRTLSEAIFLDV
						GLLLSCRTVSSCGAFILGWSVVI
00000	54050	<u> </u>	0.4020		064	LIPVISTRDDTSTTK
23892	54260	Α	24030	2	864	PSFRLAWPEGRLALLGELVCVP VVGRGMENGRILVSADRSGGHI
						CLKTLT\GKTIT\LEVEPSDTIEN
						VKAKIODKEG\IPPDOORLIFAG
						KQLEDG\RTLSDYNIQKEST\LH
			ŀ			LVL\RLRG\GMQIFP*RP*PAKTI
						TLEV\EPSSHHRKF*RARFOGKE
						GIP\PDQQRLIF\AGKQL\EDGRT
1						LSDYNIOKESTLHLVLRLRGGM
l						OIFVKTLTGKTITLEVEPSDTIEN
1						VKAKIODKEGIPPDOORLIFAG
						KQLEDGRTLSDYNIQKE\STLHL
i						VLRLRGGC*FFSHGIRSAQ
23893	54261	Α	24031	1	124	
23894	54262	Α	24032	97	296	
23895	54263	A	24033	71	269	AAGYRPCPRHPGARRTAATRR
						CHPSPAVARPAGCSGPP*CYAL
1						SPDPARTRRGDPQTPPCRLPRA
22000		<u>_</u>	21021	201		G STRLGLPKCWDYRREPPRPAKS
23896	54264	Α	24034	206	667	SFLTAQACEAQKTGSATRRKD
						CFLOPTFWLLPFLVLEKGAPSA
l .						CRGKRYVRAVWLSLSTGAILOP
						YPKLOVSNWL*CARDOEMPEG
1						CL*OSCOTLSLLVLPPKEKKDV
1						*EISPWTPAPPEHQYTVRTFQSP
23897	54265	A	24035	3	190	GTFKNDLILLISKSYY*APLPSD
				Ī		FSHSDTSS*KPSKSLKTYPSPPFF
			l			SAKTSLLQCPRNILY
23898	54266	Α	24036	197	474	SPLSHRGASELLSSLHLRSSCSS
			l			PKMAETPEAGFQIVHSASPQPG
	İ		l			TAPGNPWAAGAGG*HCEPGQL
1						AGPYRAASFFGQAWPPHPRAK
						LGVSR
23899	54267	Α	24037	164	323	GQEIKTILANMVIPRLY*KYKKI
			l			SWAWWHAPVVPATWEAEAGE
L			L			LLEPGRQRLQ

SEQ ID	SEQ ID NO:					Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540.217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		03/340,217	sequence	or pepulae sequence	decision, possible nacional insertion,
		<u> </u>				
23900	54268	Α	24038	582	1166	HCEPGQLAGHLGQPLSSGRLGF
				1		LMASHKLSNNYYRTRDGRRES
1		1			1	VPPSIIMWSQKALVSG\KPAERV
						YGDANHTVGNMCASRLGGYL
ł		1				GFSIRIVVFIWYLGLSVVVNML
ŀ						VGRQCDWQGLVYGQPELLVFS
						GNMPGEVARGKHRNEKAPLSM
		1			!	RPASQCRVFCVLKGCSERSRHK
		l				EQTPDTTIFKNCNTHCEGPWLH
23901	54269	Α	24039	1214	3878	
23902	54270	В	24040	62	1242	
23903	54271	Λ	24041	74	408	
23904	54272	A	24042	790	1877	DEIVDPGSMKWSYKSQLNYKT
l						KRPSVFVTALTHRPDILILGTLE
						SGHSRNLTCSVPWACKQGTPP
						MISWIGASVSSPGPTTARSSVLT
		l				\LLPQSPQDHGTS\LTCQCT\LPG
1		1				TGVTTTSTVRLDEAYPA\WNLT
		1				MTVFQGEGTASTTLRNGSALSV
l		1				LEGQSLHLVCAVDSNPPARLS
ł		1				WTWGSLTLSPSQSSNLGVLELP
						RVHVKDEGEFTCRAQNPLGSQ
		l				HISLSLSLQNEYTGKMRPISGV
						MLGAFGGAGATALGFLSFCIIF
		l				VGVRSCRKKTARPTMG\VGDT
						GMEDT\NAVRGSA\SQGTPELN
1						PRQKEQAPQNHGSANALANPP
į.	Ì					RPEEGEIQYASLSFHKARPQYP
1						QEQEAIGYEYSEINIPK
23905	54273	Α	24043	183	1176	VHQLVFPRQHQRHSVFPKANSS
						WIHQLFPQ\QHSYGPSAVSAGQ
						HQLMVHSLFPQATPAHASISLV
						FRQGTHQLHRVHQRRIPSAQHQ
	1	l				RHRLHQRRLPSAQHQRHRDSIS
	1					VCFPFRAHTRPHCDSIQRQHQL
						MRSIGLFSSGKTDPTVTLLKEA
	1					AAEHEICLCSRHPVNLRETKAH
1		l				LGNAAHA VFPRGNQKKPEMCP
	ł					YLPTHKLVVPTPPLIPLEAALRN
						IAHSRSIPPPKKFSPTPDTSDDYL
						LVFNSFPYSKLTTPKTGQGENL
1	1	1			1	GSTRPIPCNGPPTRGPHNGAKY
		l				PGDVSHPVGVGTLWANRLLTG
		L	L			QYEENGKSFFLLICTIATFQVNA
23906	54274	С	24044	178	342	
23907	54275	С	24045	86	283	
23908	54276	С	24046	91	177	
23909	54277	С	24047	H	169	
23910	54278	С	24048	209	673	
23911	54279	С	24049	164	208	

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23912	54280	A	24050	182	255	DAESRRSWQPLPEPLELRP\RLS D
23913	54281	А	24051	146	456	LKNHKRSEKPCPTLTDDIPPQK KCKWPVLALSADITL*KSFSWL ILAQKAPPLSTL*PPLLPAREQM SFD\LIFLYLPKSYKVAPPLSSFT ESLFGLSPPAPR
23914	54282	С	24052	90	215	
23915	54283	В	24053	1	1059	
23916	54284	c	24054	129	215	
23917	54285	С	24055	19	347	
23918	54286	C	24056	228	260	
23919	54287	В	24057	406	602	
23920	54288	C	24058	7	375	
23921	54289	A	24059	322	400	FLLSRVPSV*DPTENQTVQLTW EWLP
23922	54290	A	24060	818	1078	ILDVFSYTMKIEPSPLRSYPISPH YNL*QLPP*LDP*ESGYKTPLSA LLLTFLLYISSFASHKVSSSSVD PLPTCAYLPIGQAHVH
23923	54291	С	24061	1	411	
23924	54292	Α	24062	1	861	
23925	54293	Α	24063	425	484	
23926	54294	A	24064	1844	2097	HSAGTKLRLVNFLLTLRRLVNS VMRLKKSFMMVQRMVSIPKV WSISLGFLLSFVPSVWDPT/ENR TVQLT*HPLPEPLELGPRLSD
23927	54295	Λ	24065	729	807	LFSCSCPPKPPRP*LTAEKGGLC IFL
23928	54296	В	24066	1	1317	
23929	54297	В	24067	509	810	
23930	54298	А	24068		385	MDLPSTVRVTRSSASVMVQGG SEAIRQRQSSAAKPRRSQKESKS LGPEPQGLWEWLPEIYFSQFRR LGNPKSRHQQIQCLMGACFVV TERLSFQDSLDLAEPRHSYPLQ VKRLDLPTVLETETDLEKLDPF TTIPSPVETLQCLFVSLRIKSRVP *FWISQPPEL*EINFWQPLPEPLE LWPKAL
23931	54299	В	24069	1	774	
23932	54300	Λ	24070	350	640	SLQGCLSDNSPTFQRCQTTQGR LPWSFTLSGKSHFSGGVARACY TCQKSGHWA\RNASSPGFLLSR VPSVSDPTENRTVQLTWQPLPE PLELWPKAL
23933	54301	В	24071	85	331	
23934	54302	В	24072	1	1844	
23935	54303	В	24073	1	1169	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23936	54304	A	24074	63	220	NLTGNIWNREWYSHFDFDLLD PVGFSGSIAKMPKRKLFKCHSV *GHWRSKSE
23937	54305	В	24075	497	898	-
23938	54306	۸	24076	3	678	GGGFSDRDRTIALQPGQQIWM WGIGKQVGGCGWLRGKNRVV GAAKDQSGTGEKFYFIPKAMG AMEGSGQRRTLPDPGAIRNPS GGYSATDCEGSRPAPGEINSHA SHTKPVWWFFTQTRMKFGAVT RIGGLPWEVNPLSSCSLLRE/KS PPTTSGPQSORPKHLT-KFKS A/CYTCRKSGHWAKECPLPGFL LSRVPSVWDPTEN/KTVQLTWQ PLPEPTELLWFRLSD
23939	54307	В	24077	522	873	
23940	54308	В	24078	329	530	
23941	54309	C	24079	147	439	
23942	54310	A	24080	3	3889	
23943	54311	A	24081	2	394	
23944	54312	Α	24082	1	197	
23945	54313	A	24083	1 .	205	CLRAGPHALYLLC/NIPEFTFL\S RADF\ARSQLVDLLTDP\FQ\QEL EELLQVGAVTLILRILPCLPCLP
23946	54314	A	24084	134	182	SCFIPSSQPVAGTERSRFLGPQY FQTPHNPSGPGLGNQLMR*AF
23947	54315	A	24085	257	481	
23948	54316	A	24086	415	630	
23949	54317	Α	24087	123	340	
23950	54318	В	24088	49	270	
23951	54319	A	24089		900	RSRVRREVVALRRGLACIPSLG VWRFPSVGVLAVSDRWCLAVS DRWCLARATLSCFGCAMPMYQ TRRQENDLRTASIAV*RRKQON DH*KQRRWONIORKGPKRYTVI AGNSQSHQPMIFSMLRKLPKVT CRDVLPEIRAICIEEIGCWMQSY STSFLTDSYLKYIGWTLHDKHR EVRVKCVKALKGLYGNRDLTA RLELFTGRFKDWMVSMIVDRE YSVAVEAVRLLILILKNMEGVL MDVDCESVYPIV*ASN*GLASA VGEFLYWKLFYPECEIRTMGGR EGOROSPGAGATFFOLLL

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1	l		sequence		
		<u> </u>				
23952	54320	A	24090	1	812	MARTPGAELAVAKYPKKGSQA
		ŀ				VHRHSRKQSEPPANDIFNAAKA
	1					AKSDMQHREVRVKCVKALKG
		l				LYGNRDLTARLELFTGRFKDW
						M\VSMIMDREYSVAVEAVRLLI
						LILKNMEGVLMDVDCESVYPIV
		1		l		LFYPECEIRTMGGREOROSPGA
İ		1		l		QRTFFQLLLSFFVESKLHDHAA
		1				YLVDNLWDCAGTQLKDWEGL
1		1				TSLLLEKDQSTCHMEPGPGTFH
1		1				LLGESPGISAIVQGAIEKSGERSS
		1				PPPKRIGSMKEGPMVLPLALDT
						OCWFPVEVPLD
23953	54321	A	24091	152	1209	MPLKNVQNITHFPLEKELALPL
23,33	34321	ľ`	24071	1.52	.20)	LIFSPLHWCVSGYLSLPLPNPKP
						HFSNPLLLQVILPALTLVYFSIL
				ŀ		WTLT\HISKSDASPGECGS*MG*
						WEQRNLCPSILPLQSPLCPTEAA
		ŀ				VEFEGTRMVAFCELC\QSCLSD
1						VDTEIP/ESR*VLAOSGRKKAEY
						PCPPFPRRN*QCHYLPWETFCPF
1						KHPFFSPLQAFVLLSDLLLIFSP
		1				QMIVGGRDFLRPLVFFPE\ATLQ
ŀ						\SELASFLMDHVFIQPGRPGQW
				1		CSDSIPGAQ*GLGLGCSSKI*GR
	Ì	1				KLKSVIPFCLPCEPGDSQEDHLQ
						IERLHQRRRLLAGFCKLLLYGV
						LEMDAASDVFKHYNKVHQGPS
		<u> </u>				LGFTKPLVMPRFSAATV
23954	54322	A	24092	37	625	SLMLLTPFHFPPEECPPPIPTAH
İ						HLLPFLLSTWELFLQQLQEVVV
l		1		1		KHAEPAVLEAGAHAL\YLLCNP
1		1				EFTFFSRADFARSQL/V*DLLTD\
						RFQQ\ELEELLQGRELGAGTMG
ì		1		1		DTPNKGWGGGCKTGGMWDL
1		l				MSSGGKCYCGIFFSFSILPSQIED
						EVYNLGSHSETPLCLLQVSGFP
						PLPHPVFTGVCGVGSLCSFPV
23955	54323	Α	24093	2	198	PNDRSRVTREMSAPSRSLGRFL
		l				SVGVWHAPPSLALVPPCRCTRL
1	1	1		1		KTYHGRGAPLRVELPTCMYRL
						SNVQGRSGGPAPGVGHLQSLV
1		1	1	I		DEWLDSYKQDQ\DAGFL\ELVN
1		1	1	1		FFI*S\CGM*RNSGDYPLTAPGL
1		ı	l	1		SWKKFQGSFCEFVGTLVCRCQ
1		1	1			YILLHDDFPMDNLISLLTGFSDS
1		1	l			QVCAF\RHTSTLAAMKLMTSLV
1		1	l			RVALOLSLHODINORQYEAER
		1				NKGPGQRAPERLESLLEKHKEL
1		ı	1	1		H*YQAKNLSRARCASTCGASH
23956	54324	В	24094	211	3841	Q. Markold McAdire GASII
23730	J4344	10	24074	1-11	50	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23957	54325	Α	24095	1	1798	MSLLAFKIILVGFEGSKARVQT
		1		l		DLFYRSLVLFLLLQICFPLSCFG
						NWKGEEGGMQRWVLGKNEKI
						ASPHRLPAPTARFAVSLQLNLRI
		ł				LGAFQFGPVLFVLWRPLDRDN
	İ	ı				ASPQSEFLNSCGITAYLVGKRE
		1				GGVERGGGGRSSERCRTRLSFS
		1				PRHSHVFSTSRFCKILKKSPGAN
		1	İ	1		GGRCRARGGGLHGARAANPPT
		1				PIAEPARLPACLSRAEPEPQAAE
		1				PAPPPVGVVRSRPLSHRRHPVV
		l		ł		CPAAPIRLINKRLAHSARPAGIC
						CVSRSGLSALAAAGTASMFQLP
		l				ILNFSPQQVAGVCETLEESGDV
				1		ERLGRFLWSLPVAPAACEALN
						KNESVLRARAIVAFHGGNYREL
		l				YHILENHKFTKESHAKLQALW
			l			LEAHYQEAEKLRGRPLGPVDK
		l				YRVRKKFPLPRTIWDGEQKTHO
						FKERTRHLLREWYLQDPYPNPS
		l				KKRELAQATGLTPTQVGNWFK
		1				NRRQRDRAAAAKNRSQTSPWA
		i				EAFRFPECGSRLAKHFAEEPGLI
		ĺ				PQAQMLLTILFAFRSGRPDPTRS REHLLRRKPOADAPLEM\LOOO
		ŀ				VLSOGSGRALRAEGDGTPEVL
		l				GVATSPAASLSSKAATSAISITS
23958	54326	A	24096	3	1034	GLHGARAANPPTPIAEPARLPA
23936	34320	n	24090	ľ	1034	CLSRAEPEPQAAEPAPPPVGVV
		l				RSRPLSHRRHPVVCPAAPIRLIN
		l				KRLAHSARPAGICCVSRSGLSA
						LAAAGTASMFQLPILNFSPQQV
		l				AGVCETLEESGDVERLGRFLWS
		1				LPVAPAACEALNKNESVLRAR
		1	ļ			AIVAFHGGNYRELYHILENHKF
		l				TKESHAKLQALWLEAHYQEAE
		l	1	l		KLRGRPLGPVDKYRVRKKFPLP
		l	1			RTIWDGEQKTHCFKERTRHLLR
		I	1	l		EWYLQDPYPNPSKKRELAQAT
		l	1	l		GLTPTQVGNWFKNRRQRDRAA
		l				AAK\NRLQQQVLSQGSGRALR
		1				AEGDGTPEVLGVATSPAASLSS
		I				KAATSAISITSSDSECDI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			1	sequence		
23959	54327	A	24097	3	11024	GTRVGVAWRSVRLLLGPGAGL
23939	34327	Α.	24097	3	1024	RGGVVLKVSPSPCRGGRRVPVR
1		l				AEGARGRGPINKVPGASSGAA
1		l				VGAARLTAPLLAKAMASKLLR
						AVIL\GPPGLGKGAPVCQR1AQN
1						FGLQ\HLS\SGHFLRENIKA\STE
		l				VGEMAKQVYREKVFLVPDHVI
į.						TRLMMSELENRRGQHWLL\DG
					1	FPRTFGQAEALDK\ICEVDLVIR
		1				LNISFETLKDGLN\RRWIHP\PSG
						\RVYNL\DFNPPHVHGIDDVTGE
				i		PLVQQEDDKP\EAVAA\RLRQY
ł		l				KDVAKPVIELYKS\RGVLHQFS
1						GTET\NKIWPYVYTLFSNKITPI
				l		QSKESILTLAQWEEPGRCGSFIP
						IVWCSNWCCCPNLEA
23960	54328	Α	24098	2	178	
23961	54329	Α	24099	96	340	FGALTRIG\DLPWEINPLSSCSLL
		l		1		PEKDPPMTSGPQTDQPKEHLTN
						FKSAARRSSRTSSHRNLLHVLEI
1		1		i		WQLDQGMPAARDS
23962	54330	Α	24100	3	155	
23963	54331	Α	24101	1	162	
23964	54332	Α	24102	1	1032	
23965	54333	A	24103	167	476	TAMLLTQSLFGGLFTPTHMKFG
			ļ			AVTQIG\DLPWEINPLSSCSLLR
				l		EKDPPTTSGPQTHQPKEHLTNF
						KSDHHGCRASGNSHSGRMLLIF
						SPSRGLYACSCVKT
23966	54334	A	24104	371	772	
23967	54335	Α	24105	146	312	QKRSIPNSSPIP*SHSLLANCTK/
		l				DHHFLNGSKSFQPQLLSCKFS*
						GQHLFQGALCT
23968	54336	A	24106	76	441	
23969	54337	A	24107	1	1284	
23970	54338	Α	24108	150	371	EESRAAGIPWPSGQISGTCSKL\
						CGRRFWRNAWPLRFRHLEVLV
				i		CWRCGCGLSHSGGKELQLFSII
						VHLEGEVN
23971	54339	Α	24109	280	351	
23972	54340	A	24110	1	483	
23973	54341	A	24111	129	363	FGVVTQIG\DLPWEINPLSSCSL
			l	1		LCEKDPPTTSGPTKKHLTNFKS
				·		SKRPLFTLFSNLPHYPSTSFSFQS
						WRYTSISPFS
23974	54342	Α	24112	2	111	
23975	54343	Α	24113	3	808	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( N=Unknown,Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)  TALLLTOILFGGLFIWTRVKFG
23710	34344	,				ALTWIGDLPWEINPLSSCSLLH EKDPPTTSGPQTDQPKKHLTNF KSACFERIKACYHSPATAWPFK AYKLSLQFPHFTCPKTR
23977	54345	Α	24115	ı	753	
23978	54346	Α	24116	1	513	
23979	54347	Α	24117	625	701	IHVAHKLAVVMPGCKRGCLSL FYFQLIWSHKGWVRSGEASHR EMAGLLLLCSS*PPSPCETLPRS EPSLCDSRSAESRTKTSNLSCIP ASLQQVCGPREFIWSGNHCHPF LPITRCYGCLCYHYCMENSIL
23980	54348	A	24118	458	605	VVEEGIEDKRVYRLGTTGWIGK TIWLVRRRS*TNL*DLSGFWTG KMGEL
23981	54349	A	24119	566	769	IKEKGGLFSFSASNLWKQVTSS FLCSPLPLSLLKSSKLLANRDKY RMCGPVPANGNRTQQ*SKHVR
23982	54350	Α	24120	345	664	GEIAASQEPAGGFSGSSAQPCPL PWTGPPTTQAPGH/CNSAPSSRD NASSNGGRRQRVPTVPGGGLIL QREHILLPLGPALRAAGKEPAT AATIPESPGGDSDDSRT

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l	l	sequence		
		<u></u>			L	
23983	5435I	A	24121	2	2204	IDMIFTPGPPSTPKHKKSQKGSA
						FTFPSQQSPRNEPYVARPSTSEI
				l		EDQSMMGKFVKVERQVQDMG
1						KKLDFLVDMHMQHMERLQVQ
						VTEYYPTKGTSSPAEAEKKEDN
1						RYSDLKTIICNYSETGPPEPPYSF
1						HOVTIDKVSPYGFFAHDPVNLP
			İ			RGGPSSGKVQATPPSSATTYVE
1				l		RPTVLPILTLLDSRVSCHSQADL
1				i		QGPYSDRISPRQRRSITRDSDTP
	1			ł	•	LSLMSVNHEELERSPSGFSISOD
					İ	RDDYVFGPNGGSSWMREKRYL
					1	AEGETDTDTDPFTPSGSMPLSST
						GDGISDSTVPLFLSSEILOKOVG
						QSITSMLGFLSRGPSMKLCMGL
						ACVLSLWNTVSGIKGEAKKEK
		l				GMTFLPTTDSKKFFSLLSVTSYS
1		ı	1			SFAFHKFSVAVYNISNLKTVDP
						AKFPTRYCYCLNNRTNDLSDFT
						ALLVDIIGNSTSYLTEIFKSTSIL
						SVNQSNESDCIFICVMTGKSGR
						NLSDFWEIEEKYPIINYTFTSGL
						SGVLALLLTQSLFGGLFTRTRM
		1			1	KFGAVTRIG\DLPWEINP/LSSCS
		l		l		LLHEKDPPTTSGPQTDQPKKHL
					1	TNFKSAARPTFLGQGQVPLNPF
						SFTLSGKSCFPRRQEPPNRLFPH
1			1			PNLLSLCPNPLFLCPNPFSTFLE
		l			:	GKNHHPHPFSVSQVFSLGLPPS
		ł				LWEHSPMAHRLRVOPROOPPDI
		1				HKQVISLQSGHSLLVKSQSWSQ
		l				DPIKFQLPHHHKNTASVCINPSV
23984	54352	A	24122	2102	2529	ETTMSFIFSLLYPFRMHLSFLIVS
23704	34352	ľ	24122	2102	2027	PALRSISRRALAAFPRGRSLDLQ
		1		1	1	PAMPEPPTPS/CGLLCVPS\PPMS
		1	1	l	1	
			I			AAPCSKVPQSHLFAFMSCNTAK VCSFTAEASETTSPPGGTNNSR
						RATLRAVTLTRRSATLTRRSAA
		L				SHRSSTV
23985	54353	Α	24123	299	567	TAMLLTQSLFGGLFTQTRMKF
		l		l		GAVTQI\ADLPWEINPLSSCSLL
1		1		1	1	HEKDPPTSSGPQTDQPKEHLTN
		1				FKSGETKEMHFIRGPKTPAPVM
23986	54354	Α	24124	684	822	KVKRQSYSQNTFYYPICSRH*IK
1						LOKLNSGPOTPOODLINLAIKV
		-	<b></b>		<del></del>	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide scquence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \*=possible nucleotide insertion)
23987	54355	A	24125	1163	1826	TVLLLTQSLFGGLFWTRMKFG AVTRINGBLYPWINPLSSCSLL HEKDPPTTSGPQTNQFKELLTN FKSYPTTSVAESFQSSFSTDPSD PSSPPQAARPQAKFGPNSSSASA PRPYNPFITSPHTWSGLOFHSA TSPPPPAQOFLKKVAQVKGT KWLKTDAARSPQKPTGTTTDAL DNSHSGGFIMDRFLLVKSPKGF LRLLYFSKFISTILINLQE
23988	54356	A	24126	636	922	TATLLTQSLFGGLFTRTRMKFG AVTRIGDLPWEINPLSSCSLLRE KDPPTTSGPQTHQPKEHLTNFK SDKGDMFYPWTQNSSAGHGLG RQPSLGV
23989	54357	Α	24127	728	1013	MKFGAVTRI/RGDLPWEINPLSS CVLC/CREKDPPTTSGPQTDQPK EHLTNFKSGKRPAFYSLLQPPSL SLNLFLLSILAPHFNLSLLLISIPF IFW
23990	54358	В	24128	I	3993	
23991	54359	В	24129	105	420	
23992	54360	В	24130	45	2866	
23993	54361	В	24131	I	1680	
23994	54362	Α	24132	1257	1403	
23995	54363	Α	24133	1	2130	
23996	54364	Α	24134	1	725	
23997	54365	A	24135	2825	4123	RAADGVITLEDNDPCVKDDSSR FPYEDRLDLVLKGTADIPRLTV HRGSEYIISRATFFCYEIKEQSVI NHCYTEIDLKIFRGYLAPALGV HREFYGTEPCRVTAQYNQDM RYWLETPTISAPPIELVEIERLRY QEMPIYASRWHIFREPDTALP RNDTPOMVKLFMDREQVVKDI GYIELKVVEDQRTRAVMIKFRA FVKEGA VIVIRFDNKEVAFAQM SGNLTVPRSTSDYKARDIFFQY REITHISTIGHGVANNHQIRLRL ELAGIVPLNQIAALLKVYPLS KLCNSPKKCQLESVFRLTQWV VKKPISNRQSGLGGIVLPGLFRV LSPGNLSPIKGTVILPNEVQGF TGGNLSPIKGTVILPNEVQGF TGGNLSPIKGTVILPNEVQGF TGGNLSPIKGTVILPNEVQGF TGGVSNAGLADHTRASSPERA GVYPHMERCAHRASRYY*KW HPDSVAADHLLG

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nuelcotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /possible nuelcotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23998	54366	A	24136	137	1199	ISSGGPRELGLKMIIVPGSQLSQ
		1		\		LPPGLDSTEFAAIVGNLLDNAF
						EASLRSDEGNKIVELFLSDEGD
		1				DVVIEVADQGCGVPESLRDKIF
						EQGVSTRADEPGEHGIGLYLIA
		1				SYVTRCGGGITLEDNDPCAPLT
		1				LLIVEDETPLAEMHAEYIRHIPG
		l				FSQILLAGNLAQARMMIERFKP
						GLILLDNYLPDGRGINLLHELV
		1				QAHYPGDVVFTTAASDMETVS
	ľ	1				EAVRCGVFDYLIKPIAYERLGQ
		1				TLTGFP\QRKHMLKSIESATQKQ
						IDEMFNAYARGEPKDELPTGID
İ		l				PLTLNA VRKLFKEPGVQHTAET
	1	l		1		VAQALTISRTTARRYLEYCASR
		1				HLIIAEIVHGKVGRPQRIYHSG
23999	54367	A	24137	2	243	RPGRGLGQRGIWEPAA\LLDSV
						AQQQGAWLPHVADFRWRVDV
						AISTSALARSLQPSVLMQLKLS
						DGSADLEKRCERRLQD
24000	54368	A	24138	1	1483	MKLMETLNQCINAGHEMTKAI
		1				AIAQFNDDSPEARKITRRWRIG
						EAADLVGVSSQAIRDAEKAGR
						LPHPDMEIRGRVEQRVGYTIEQI
1					l	NHMRDVFGTRLRRAEDVFPPVI
		1			i	GVAAHKGGVYKTSVSVHLAQ
ŀ		ı				DLALKGLRVLLVEGNDPQGTA
	1	l				SMYHGWVPDLHIHAEDTLLPF
		l				YLGEKDDVTYAIKPTCWPGLDI
		l		1	ŀ	IPSCLALHRIETELMGKFDEGKL
1		1				PTDPHLMLRLAIETVAHDYDVI
	ł	1		1		VIDSAPNLGIGTINVVCAADVLI
		1		1		VPTPAELFDYTSALQFFDMLRD
		1		l		LLKNVDLKGFEPDSDLRYLQSC
i		1				EWGDTGIFLKWSCIPAEKGLQT
		1				SGAWVEAADIPLLPGSPRRLSP
				1	1	QAGSRGGQGPKHGQQCLKMP
l		1			I	GPRAPGLQGGSNRDP/AP/GPAV
l	1	1			1	ERAPAAALSSTTIWMPMSLCP
					l	WRPVLAACTSMTVRGRSTMYL
					1	PTTTGNAGCTWPKASLATRWL
		1			1	SSPMGRQARSLMLGAQVTLSW
	1	L.	l			DRSMPWRRRGRSSGRNLSTI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24001	54369	Α	24139	727	1498	GICIWDRPPGLADQRRSHSHGW
					1	WGELATPLPWHH/HLSDSHSG\
		ĺ				RVSFLGAQLPPEVAAMARLLG
l						DLDRSTFRKLL\KFVVSSLQGED
İ						CREAAAAVLGSAPTCRR\AAGC
					1	
						LLAGMHTLLQQALRLPPTSLK
					l	PDTFRDPAPGALHPPRPGRGL\G
		1				QRGIWE\PSGPSFDSVAQQARV
		1			'	AWLPHVA\DFR\WRVDVTIST\S
	l				i	AL\ARSLQ\PSVLM\QLKAFQNG
		1			İ	SSIPAFEVPPQPKF\QEL\RYSVA\
		1				LV\LKEMADLEKK\CE\RNLQD
24002	54370	Α	24140	1	312	
24003	54371	Α	24141	2	524	
24004	54372	A	24142	144	494	AGSCPFAAGPGPLGSGRCFVLIR
	ŀ					FILI\ONRAGKTRLAK W\YMOFD
		ŀ	·			DDEKOKLI\EEVHAVVTVRDAK
			ŀ			HTNFVGVSGTFKIILPPLCWPLL
	l		ĺ			LASVWMFNDNKPGLYLGGPFH
			l			NFRGRS
		ļ			700	NEKGKS
24005	54373	Α	24143	1	428	
24006	54374	A	24144	108	422	
24007	54375	A	24145	3	387	
24008	54376	A	24146	3	337	TRLMLLDRSGQGKVYGLVGR*
		ł				RFQQMDVLEGLNLLITI*RNRN
i						KLRVYYLTWLRIKIMDNDPKLE
			ŀ			KKHDWTTAGYMEGCGHYRVG
İ		1				KYERIKFLVIALKSSAEVYAWA
	ļ		ŀ			PKPY
24009	54377	Α	24147	1	340	ERTRMNKQHNSTLAKSRPGNT
	1	1	İ		1	GPEPLMPQASPGPPGPLSQTPP
	1		i			MHRPVEPOEGPHKSLVAHRVP
	l		l			LKPYAAPVPRS\QSLQDQPTRN
	1				1	LAAFPASHDPDPAIPAPTATPSA
1	1	ĺ				RGAV
24010	54378	A	24148	335	869	ATPTLILSOKSHFWPOPLSSVLK
24010	34378	l <sup>A</sup>	24148	1333	009	
	1	1			l	LTSGPEISLQSSPSPICRMKKRR
	1	1			Į.	LSPLASGLES*VKSGEWCEVLS
1	1	1	4	l	I	RAPLFP/HNPSQAAP/EWEMQG
1	1	1			l	TRPSIPHPLGLAGLPTQLQQGRL
1	1	1			l	WGYRNPASPFRTRVAARDCAG
1	1	1	1	1		KQVRTGPDARLEGSRPGVRAG
1	1	1		1		LGWGRGYSERVQTCAKSSVSPS
24011	54379	В	24149	158	5640	
24012	54380	A	24150	ī	1854	
<u> </u>						

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	
i	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24013	5438I	Ā	24151	1	1510	MPCGKQGNLQVPGSKVLPGLG
12.013		l''		ľ		EGCKEMWLRKVIYGGEVWGK
1		1		l		SPEPEFPSLVNLCQSWKINNLM
1		l	ŀ			STVHSDEAGMLSYFLFEELMRC
						DKDSMPDGNLSEEEKLFLSYFP
		1				LHKFELEONIKELNTLADOVDT
1		1	ŀ		i	THELLTKTSLVASSSGAVSGVM
İ					l	NILGLALAPVTAGGSLMLSATG
1						TGLGAAAAITNIVTNVLENRSN
1					l	SAARDKASRLGPLTTSHEAFGG
		1				
						INWSEIEAAGFCVNKWVKAIQG
		1				IKDLHAYQMAKSNSGFMAMV
			1		l	KNFVAKRHIPFWTARGVQRAF
i						EGTTLAMTNGAWVMGAAGAG
						FLLMKDMSSFLQSWKHLE\DG
					1	ARTETAEELRALAKE\LEQELD
					ľ	RLTQHHRHLPQKASQTCSSSRG
						RAVRGSRVVKPEGCLATSQPH
1						QHQQERGDRPREDTDNEKRGG
}		ŀ			l	EKWSDFGSVLKGEPTGFVHGL
1		l				DLWCEGKKSQDNGGLWPEQL
						AGWSCWHLKWTGGSRSGLPVP
						VLDAACTGTLSPDLTSWRYHPS
		_				SIPLPKQPLQHGF
24014	54382	A	24152	3	515	VGTQAGAAPGPVQFLSSAWVA
1						L\HETELPPLTAREAKQIRRE\IS
1						R\SSKWVDMLGDWEKYKSSRK
		1		1		LIDRAYKGMPMNIRGPMWSVL
i						LNTEEMKLKNPGR\YQVRSARA
i		1				QPTGQAVSGAQVSSWRERQDH
1						PGELGVKIMKEKGKRSSEHIQQ
						MDLDVSGTLRRHIFFRDRYGT
24015	54383	A	24153	1	1317	
24016	54384	Α	24154	1640	4841	TITGSSPLAGGHHSTCRGRMDM
			1			VENADSLQAQERKDILMKYDK
		1				GHRAGLPEDKGPEPVGINSSIDR
			1			FGILHETELPPVTAREAKKIRRE
		1	1			MTRTSKWMEMLGEWETYKHS
						SKLIDRVYKGIPMNIRGPVWSV
	1		1		1	LLNIQEIKLKNPGRYQIMKERG
		1	1	1	1	KRSSEHIHHIDLDVRTTLRNHV
				1		FFRDRYGAKQRELFYILLAYSE
1					1	YNPEVGYCRDLSHITALFLLYL
1						PEEDAFWALVQLLASERHSLPG
1		1	1		1	FHSPNGGTVQGLQD
24017	54385	A	24155	1640	4056	, = ,=
	12					·

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
24018	54386	Α	24156	479	3902	GQVDEDV\$VKMDTVPGVNLSC
				l		ILNEMRDQDKKLVEKSYKDAK
						GCFFSMGEPSLIDPCLEVCYGT
						QLAQLQGLIRSMEQQLCELCCD
			1			AEHQDHEHQVLLDVKTRLEQEI
						ATYSRLLEVEDAQQGPIVMRA
						MTSECHWCLAPIRSPDCVCVLR
						QLHTRSGIHKRFCGLKGFLPER
						HNPGQLIHQNQSGCCSRMYPFD
						RKLGISVALNIPGGRSYRDLWC
						LTAVHIHIPGIDHHRLFTLGRM
						DMVENADSLQAQERKD
24019	54387	A	24157	780	983	QHQSLIGHKYQRSHRSL*AEPE*
						LECHQPDLAEPADGPPLTTRYG
						IVNAEMGLEQLLPKGPSLKEAK
24020	54388	Α	24158	773	2016	DWPTLRENRKEPTLKYWGLVP
		l				LISGMPTWFFFFLSACGNLIPSG
		l				GIRVPMENML/RVFRAVEEHCP
			ĺ			WFPEKGTLSVEQWDSVGAKFQ
			ŀ			ELVPTGNYVPITVWGDWALLL
						QFSESGDPLPLPQLSSPTGPSLS
						DQPLPSPTPPPPDDVENSISNSG
		l	l			DFGLTLPPGDLILFPEEPLLAAS
						AAPNRTALGHIYANSSLFKPLQ
		ŀ				HLPLESANGSGAKLQFTYNSAG
		1				PSPSSAAPRPPVVSVPQPVTLPS
						TQAASLYPSSHMDTSNHQCTSA
						SSAPPMPLSHTLIPSDETTCNVF
						NYPLKKPTGRELREAKNKGSQP
	1	ĺ				KVHEKTNPGNIRCISTEERPHEN
	l	l				KGFNHLLEWLAELKTTIHLLDC
	ĺ					WFIAKDVNREQPDGRDAGDKR
						LCVKFLREIAANTYKTEVGAAK
						VMEILLHSHGATTTIAN
24021	54389	A	24159	365	675	
24022	54390	A	24160	I	1191	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
24023	54391	IA	24161	1363	1434	QQGVCSGEKGGHWAPROVGV
						YLLPGRVGCVSSRVSPSFPGDG
1		1		1		LDSGLARRGSAVSALASGLVEE
		1		1		PMLGPPFHPTPRFKAVSAKSKE
						DLVSQGFTEFTIEDFHNTFMDLI
		1				EQVEKOTSVADMLASFNDOST
1		1				SDYLVVYLRLLTSG\YLQRESQ
		1				VLRALHRRVDGTVORSSAKOE
		1			l	VEPMCKESDHHSHHCAGPGPS
		1				ACPSRVGVHGTAARAAPPIRTS
		1				F/H*GVPSPKVYLLYRPG\HYDI
		i				LYKLGLGSSPLGCPGCPLLARA
		i				LGHCYRGFSVVVNGPISPPSSCH
		l				MTPPPCFIKGGAGGEPCVRVPA
		l				LLPAWLLCLLPPPPRWVPLLFT
		l				YLLLSFPNRSRFEGPGLLEAPPA
		1				SLGSASFPS
24024	54392	Ā	24162	2	888	VFRSRHOGCLKMAAEEPOOON
24024	34332	^	24102	-	800	OYPLGSDSEGVNCLAYDEAIM
						AQQDRIQQEIAVQNPLVSERLE
						LSVLYKEYAEDDNIYOOKIO\D
						LHKKYFVHPQDQGLTGTCFYS
		1	İ			GFRDSPTWEGTCWIDSKELORF
ŀ		1	İ			QGLVS/SQRSKGKTLVSQGLSL
		l				EFHNFEGFSTTTFI/VDLIEQVEK
		1				ADLCRRPCLASFNDQSTSD\YL
						VVYLRLLT/SRAYLORESKFFEH
		1				FI\EGGRTVKE\FCQQE\VEPMW
		1				QGEATTSTSLALAQALSVSIQV
l		1			]	EYMDRGEGGTTNPHIFPEGSEP
		1				KVYLLYRPGHYDILYK
24025	54393	A	24163	3	819	EGATMAAPPOPVTHLIFDMDG
i		1			1 '	LLLDTERLYSVVFQDI\CHRYD
1		l				KKYSWDVKSLVMGKKALEAA
		l		l		QIIIDVLQLPMSKEELVEESQTK
		1		l	ŀ	LKEVFPTAALMPGAEKLIIHLR
						KHGIPFALATSSGSASFDMKTS
		1				RHKEFFSLFSHIVLGDDPEVQH
		1				GKPDPDIFLACAKRFSPPPAME
			l			KCLVFEDAPNGVEAALAAGMQ
1		1	l	l	l	VVMVPDGNLSRDLTTKATLVL
1		1	l	l		NSLQDFQPELFGLPSYELEGRA
				1		SVFRPQPTLMVHTAGGKGKEIS
						NSSIPTCAVILAS
24026	54394	В	24164	115	594	
24027	54395	A	24165	4	160	SIIFPAIPVSLK*IKPNCPEPPPDH
						LLQAAWGCVSLGQGLSYSALN
		1	l	I		TLLYFT

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24028	54396	A	24166	88	2280	VPLYDRMGGETGLLIPSPERSL HAATDAHSCKNRHSLTGKTDS LWQQDTKLQTRPKATQGKGES GGCGPHRPLLLLRTKSTPESKS AHDPPVAPAQAGASGCGRGAA QKGFGAKAVDGGGRDRTPG VPAAGPPTSSGRGDRGPSSSRR
						TRSQTPESPTGRPGSRQFQAAPP PSHHSRPAHAPFPGFQMYRGPL YSSSPAPATPNTSAYESPRFCST SYISSSPRFCPTSYICSSPRFCSTS YICSSPRFCSTSY
24029	54397	A	24167	2	352	WRTLIVLGMSCLGSYPK*MHLT HLIHCS*EARSLVTLHIPLTICG EPRNIIKLVGCP*IQWKK**KKI MNSRILSPSFRSRY*TSNLLGLP *LRGLSPVERELKLWTNRHKLL
24030	54398	А	24168	2	252	NKSSVFATSCSLDPISIMSLM*W TSVISCRSKK*SRSLRIRL*HKA KELIYPLPVRESMWGYCQLLS MSMPIMHAGTGEMTTG
24031	54399	A	24169	1505	1680	RTSSLVTL*TLHLEPAGT*VLTP GSTTGITVSGTVLFLILLSCFLSS TPFRLKLACLS
24032	54400	Α	24170	681	932	
24033	54401	А	24171	1740	1950	HHHHLLQMSWSAGMQAWHW PSFYNVVTGKTLA/LPNLIALQH IPLSPAGVIAKRPAPIALPNSCA ALLSSQ
24034	54402	A	24172		652	GKLITEARTRCCMQLTSDRINGH TALNHTTCSHICVSGMAGRL GPITVLSGDIGPKGMRALTMW HTLARPTGDLEAPRHGPRRIT ATOPPHAPPATSRINRAAHPHN HTPPHPQTAHPTATHSHLTYR TPRTPTATRTPLHPHSHQVPY QPTQRSPYLHPPPAPPHISLHPS TITRATVNAPDTRFITYARESVPF GRILINLGRQRGMELANILQVH GP
24035	54403	A	24173	267	400	RFLHCGNLAGLLVGFLSFKLGE YEGLGPLAALTAGKGLFFLPLE SRLRLHRRPKPRSPASFG*VSRH THAIHFN
24036	54404	Α	24174	422	452	
24037	54405	В	24175	131 .	1047	
24038	54406	Α	24176	2	199	LGLR/CARLEARARPPWATVRE KLWLFGHPTHMRCSWRGLFQD DLGLQKPLSGGSSSAYRGCENL NS

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod		location of first		*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24039	54407	A	24177	440	1746	RNLOCVFOGRIKEGIPVLALS/L
					1	RPSP/RVLHTATKQKNSCQHLE
						EDV D\QTKWRTDTPSMDKILME
		l		1	1	EVKLEEQLKEAVEEDKQALAD
		1		1		TEGSEQSSQKLVEEGNMYSIQG
		l		1		FCKDSLEVADVLEKATQCVPER
		1			1	EIKDNNPHLKNLCETLTMSEEL
		l				SQMVDCCRLPQDVSVLGVALH
						WVTPQMLCVPVNEDIPEVSDT
		l				VLKAITAPPRVPIGRTSGDVTGT
		i				GVRRRRVSSAMSQHRAFRDVT
	}					VTVQCLELGCFLLRVELASQFL
	1	1				KRPRSPESNEKVPEIEVTVEASA
	1	1				KMSHPTLYDATRSPWGMFQSQ
		1				SVLLSCAHHSLLRGATLDRCGS
						IDESEEWYGMDTFECTGVGVSI
	ĺ					RRATEPERSSGSAAALDSHRSM
						HPIVNHACEGSTLRCPYENLMP
		l		l		DALSLSPILSSRSGDASPQAEILF
		l	1	l		VVGPFRMRRCLTVDPPGLDVT
24040	54408	Α	24178	391	656	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24041	54409	A	24179	1	2150	MEDSTSPKQEKENQEELGETRR PWEGKTAASPQYSEPESSEPLE AKQGPETGRQSRSSRPWSPQSR
					1	AKTPLGGPAGPETSSPAPVSPRE
		ł				PSSSPSPLAPARQDLAAPPQSDR
		ļ.				TTSVIPEAGTPYPDPLEQSSDKR
		1				ESTPHHTSQSEGNTFQQSQQPK
l		l				PHLCGRRDVSYNNAKOKELRF
		l				DVFQEEDSNSDYDLQQPAPGGS
		1		1		EVAPSMLEITIQNAKAYLLKTSS
						NSGFNLYDHLSNMLTKILNERP
						ENAVDIFENISQDVKMAHFSKK
		1				FDALQNENELLPTYEIAEKQKA
		l				LFLQGHLEGVDQELEDEIAENA
ļ		l	l		Į.	LPNVMESAFYFEQAGVGLGTD
1		l				ETYRIFLALKQLTDTHPIQRCRF
		1				WGKILGLEMNYIVAEVEFREGE
1		1				DEEEVEEEDVAEERDNGESEAH
				-		EDEEDELPKSFYKAPQAIPKEES
		1				RTGANKYVYFVCNEPGRPWVK
		1				LPPVIPAQIVIARKIKKFFTGRLD
		l				APIISYPPFPGNESNYLRAQIARI
		1		l		SAGTHVSPLGFYQFGEEEGEEE
		1		l		EEAEGGRNSFEENPDFEGIQVID
		1				LVESLSNWVHHVQHILSQGRC
						NWFNSIQKNEEEEEEEDEEKDD
1			ļ.			SDYIEQEVGLPLLTPISEDLEIQN
l		1	l			IPPWTTRLSSNLIPQYAIAVLQS
		1				NLWPGAYAFSNGKKFENFYIG
		l				WGHKYSPDNYTPPVP\PPVYHE
						YPSVPEMAEMDDPSVEEEQAF
						RAAQEAVLLAAENEESEEDEDE
24042	54410	В	24180	1	489	
24043	54411	Α	24181	39	423	RSPTPDAASRSQCSSRRRRGVS
						KEVFVSMARTKADWPANRTG
						G*STPRKPTGPTKSPLRKSAP/A
		l			1	PTGGGERNPHRLQGLVTVAAP*
						NLDGYQKVHSNFLIRKLSLPSV
						WVREIASGLLKQNLRFQERQ
24044	54412	Α	24182	3	384	
24045	54413	A	24183	1	434	MLQNVTPARAEWQRKTALQG
		l			1	VEDPVPARPPISSPEDLLRSGEC
1						STAPTQRQVPQPVHE/SHRPTQ
1		1	I	I	1	DSPGARRKPTSERGHTEQRAAP
1		1		I	i	DREARPATRGWGLRPGGQWLG
			1			GSEHAAAAPGPSDHPRGRCRH
		L_	L	L	L	HTILQRPRLLSGKTES

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24046	54414	A	24184	15	286	LPSRGAWLGTCSPPCLSLPPPP WAPVRPEPPR*APPPAPRRPVPS TTQGSLGPHC*PLE*HSPRRSA ASLLKPARPRAHQEERTTPDAP P*EL*HSP*RSAASLLKPARPRA HQEERTTPDAPP
24047	54415	A	24185	2	214	CLSLPPPPWDPVRPEPPQ*VPSP APQHPVPSTTQGLRSVGARRGT GRQLHLQSRCGIHWVKPAGLL SLGL
24048	54416	A	24186	308	311	LPSHGAGLGTCSPPCLSLPAAP WAP/RVARASPTSAAPTPQHPV PSTTQGPLGSPSTLRKL*SFALC NKSCCCSLFGSTLPL*AVTLTA KVCSFTPKA\ARP
24049	54417	Α	24187	1	669	IMMPEGSGISLSGCDLGTALQLL SQAIVECLKLFQMLNRSTILGSG GQQPSSHSSTKLRPDSGAQLAS PSGSCTGAAGGAACQSRTVCSH SSALGWSMGLGAVEGGVVSLV GEAQAAQEPMEGVGSGMAG CRSRALPHGKAAKARRAIEHST GGLALLGDPVHSPQPLARVLSP SLPRASRAGWPAPSACPPSPHP PGTPAGPQAPHAAPVPARASPS TPPCKL
24050	54418	A	24188	157	395	RCSSCQREMTPYGK*NVQKRL KTTKNGSM*LKSHILSTPEQ\PA SPTSS/PPLLQPAAPAPPLMPRPS PPPGISVA*YPGM
24051	54419	Α	24189	2	287	CLSLPPPPWAPVRPKPPR*ATPP APRRPVPSTTQGLRSAGARGT GRQLHLQPLCGIHWVKPAGLLS LVGTWRTFMSSSGIVNTPISTLC LAQGL
24052	54420	A	24190	500	724	
24053	54421	В	24191	1	2154	
24054	54422	А	24192	688	1215	SRLAPALLSPWVVDGTGRCGA VGGAHWGGLGRTGAHGGGR LRHGGLRVPSPAPREGS*GPPR YRAQRWWAGT/DWRTQYTLRS CWFGTLPSK*FEMTPYG*NV QKRLKTTKNGSM*LKSHILSTP EQWASPTSS/PPLLQPAAPAPPL MPRPSPPPGISVA*YPGMPWTC PFYWQ

SEQID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
		ļ		1000	<u> </u>	LIVER THE PERSON NAMED AND ADDRESS OF THE PERSON NAMED AND ADD
24055	54423	A	24193	308	1131	VLQLLRRRTWSCSLRAWSWDL
		ı			i	IVPPGGLVVSLASGVKLQTFAV
						SVTALKEARLDLFVPPGGLVVS
						LASGVKLQTFAVSVTAHKSSV
		1				DPKNSGAQLASPSGSRTETAGG
	i	l				AACQSRAVRSHSSALGWSMGL
		i				GAIEQGAALVRGGSGRTG\PME
						WVGGSGMAGCRSRALPRGKA
l						AKAQREIEHSAGGPALLGDSVH
l						PPQPLARVLSPPLPGASRACWL
					ł	LRVRGPPSPRPPGTPAGPQAPH
		1				AAPVPARASPSTPPLVHLTNDV
	1				1	LRVKVKWAFFFTLHK
24056	54424	Α	24194	192	818	SWCLSRAVVGDSMTSEVIED\E
						KQF\YSKA\KTYWKQIPPTVDR
l						HAWGEGPNKTGTSCALDCGAG
ŀ						IGRITKRLLLPLFREVDMVDITE
						DFLVQAKTYLGEEGKRVRNYF
ŀ		1				CCGLQDFTP\EPDSYDVIW\IOW
ĺ						VIGHLT\DQHLA/ESFLRPCKGQ
						ASAPNGIIVIK\DNIA\QEGVILD
	1					D\VDSSVCRDLDVVRRII\CSAG
	1	1				LSLLAEERQENLPD
24057	54425	A	24195	h	233	FFFFETESRCVARLECSGAISAH
24037	34423	'`	24173	1	233	CNLCLPGASNSPASASO/CSWD
			1			YRCVPSRLANFLYFSTDGISPC
						WPRMVSNLLTS
24058	54426	A	24196	240	476	LLPQYRRVLLIYSGIQLLPGLGL
24036	34420	^	24190	240	470	RRRMCPGIYSFLLDFLVYLRRG
1						VYSIL*W*FVFVVLVPWFSAPS
						GHLRSSLCCLF
24059	54427	A	24197	288	677	NLPEDVVPDDKPHQPLLRAV\C
24039	34427	^	24197	200	077	LFVISKDLFLESPFESPIWRMYP
1						NSTVHIQRLELALLC/CHCPRAL
ł						PASAHSCHSLERGLPICKE**CM
		İ				PCSEYTIASQRPAGGAPAILPEC
1						DSRRRTTQATSEPGSHI
0.1000	54420	١.	04100	00	200	
24060	54428	Α	24198	99	399	LLCHGFEHPPLAQRSLLLPIV*S
						LLLSTRQSHSPSSFVPLLARSCG
						SWVSPAEAAEPQILQKGKCSCL
		1				IVPLEALCQRGTRLSLWLPLTFF
		_				PSFQLWPV
24061	54429	Α	24199	1	586	LSFFKVFSFFAIGSNILFSSEKFII
l		1	1		l	TDHLKPSSPTCQSHSLSSFVPLL
						ARVAFRWGFGVDVLSFVSFPSN
		1			1	SQDPQLQVCGVCWRSTPDRLP
l		1			l	GYHQRGSGGCRTADIIEPQMLR
					[	LIVRLEVLSQRSTRPQRCQ\SA\L
					1	LGGASQLGCC/GVRDLLEEAVC
		1			1	PFSDL/KLHAGKTTTLFKAVRQ/
1		1			1	SHLSLQRLLLSFVCLCPT
		_			•	· - ·

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24062	54430	A	24200	II	822	
24063	54431	A	24201	1428	1801	RSAGIDPE*FFSG/PRRIHTASCL
24003	54451	ľ	24201	1420	1.001	PSQRSSNRACSSSVTRIVSILSRF
						IGIITPMNRNPRFWRNSTSWTR
					i	MNRQTSVNRFTTTLMSFTAAA
		1			i	SRVSVMTVKTSDTCSSRRRSQL
						VKYHLSISTQVL
24064	54432	В	24202	1	2040	THE THE STATE OF T
24065	54433	A	24203	176	564	AWEPKWYSTTATRTHFWEMRS
		-		1		RSIMSTTKAFSTSNSASDSLEEP
		ı				SSRKTRSSSLLQ*KSCNGGSEM
						SSTYRVVGLTGASLSASALNSLI
						DSRSNR*V/FSRNTLANITRRNO
						LKMLPAPDPDGFSSGSPFQ
24066	54434	Α	24204	I	774	
24067	54435	Α	24205	1	1807	MPLRAFNPVSSFRWARGMTIV
		1				AALMTVFFIMQLVGQVPAALW
		1		1		VIFGEDRFRWSATMIGLSLAVF
		ĺ		1		GILHALAQAFVTGPATKRFGEK
		l			i	QAIIAGMAADALGYVLLAFAT
		1		1	ŀ	RGWMAFPIMILLASGGIGMPAL
		1				QAMLSRQPNFDHWTADRHGD
		l				LCRLGEHMERVGMDCRRRPIPC
ļ		l				LPPRVASRCMEPGHLDLNGSRR
		l				HLANGFTTPRIGANQFLRRTVN
1		l				AQTNPWQNISIASAISSSRTRRIS
		l				GSVGSWPRSGNAEVSALHHYV
		l		ļ		PDLHRRMLLATLWNTYICINEA
		l				LALTLSDFSLVPPHPYRQLFTLT
		l		ł		TFHNRACSSSVTRIVSILSRFIGII
		l		1		TPMNRNPPYTEASVTKQEKTAL
		1				NMARFIRSQTLTLLEKLNELDA
		1				DEQADICESLHDHADELYRSCL
		l				ARFGDDGNSYDHDYAKLACLQ VEGRGGGTFAHELLAVEYAGW
		l				ISPAFRLKVMHWLMNRMLDAR
		l				NRDRYKQWKQDFFHDSYNRQS
		l				AGHILSQCANLAATTSEYFIHKF
		l				HRLIAAETGYSQSTVVRAFREA
		l				VNKGILSVEIVIGDHRERRADL
		ı	1			YRFTPSFLVFAQQAKNALIESKS
		I	I			KI\SSAATKVRLLAPISPTSPMG
						KIGLATSGS
24068	54436	Α	24206	291	851	
24069	54437	Α	24207	2	694	
24070	54438	Α	24208	2	331	HLSTIEPEDYNLLFRRVSLKLSV
			1			WLPTWQSKSNGWSKKCLWMH
1		1				LKSNPLNALLPPAGF*SATMPK
		1	l			CSAWAKTISSPGCVITEF*SQPV
		_				NAGTSPNKNTYLVGISPLKKP
24071	54439	Α	24209	1	4455	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24072	54440	A	24210	698	1088	LTSKFIAAVCMRRISLCGKGSA
						QNTDSGIAATPCSCVSD*QNSTS
ŀ	1	l				SRIRM*PWMCPCPMKRKSAT*N
		l		i		ITLRC*RAAPPISSRVRK*/MSM
l		l				Y/NHLQSAECGGHRPFPVPQEE
						SILLLHLIANRSSACHSIR
24073	54441	Α	24211	1	1902	
24074	54442	A	24212	1	2898	
24075	54443	A	24213	1	2604	
24076	54444	В	24214	1	1254	
24077	54445	A	24215	1	2946	
24078	54446	A	24216	698	1285	RFWRNSTSWTRMNRQTSVNRF
						TTTLMSFTAAASRVSVMTETA
		l				MTMITPSLHACRSKAVAAALLP
	ŀ	1				MNCSLWSTQAG/CSP/DVSAEG
		1				KPDIYRLSNRKITTCYSAES/LL
						KLSVWLPTWQSKSNGWSKKCL
						WMHLKSNSLNALLPPAGF*SAT
ł						MPKCSAWAKTISSPGCVITEF*S
		l				QPVNAGTSPNKNTYLVGISPLK
		l				KP
24079	54447	Α	24217	1	1986	
24080	54448	A	24218	1	2262	
24081	54449	В	24219	1	1014	
24082	54450	Α	24220	1	1155	
24083	54451	A	24221	5	386	
24084	54452	Α	24222	1922	2146	SSSAGISCWSSSPTPGSIPSRVGS
						AAA*GWTTSRSSTGSANPSASR
						KPAAAIRASMPPNCRSGEARWP
į						LWSGSL
24085	54453	Α	24223	1091	1435	
24086	54454	A	24224	1	2919	
24087	54455	Α	24225	1324	1502	
24088	54456	Α	24226	1	2901	
24089	54457	Α	24227	5	253	
24090	54458	A	24228	1227	3031	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24001	E4450	A	24220	1	1940	MNROTSVNRFTTTLMSFTAAA
24091	54459	A	24229	1	1940	
		1				SRVSVMTVKTSDTCSSRRRSQL
						VCKRMPGADKPVEGRGGGTFA
		1				HELLAVEYAGWISPAFRLKVN
		1				QTFIDYRTGRLQPAIPQSLPEAL
1		1				RLAADLAEQKQRLEQKMLMD
		1				APKVEFAERVATASGVLIGNYA
		1		1		KVLGLGQNYLFTWLRDNGILIA
						TGERRNVPKQEYISRGYFTLKE
		i				TVIDTSNGSRISFTTRITGKDVV
		1				RTTSIVMLAKVTFLSCITMSDFT
		l				FSGYELACFVTHSGLSRSAGHIL
		l				SQCANLAATTSEYFIHKPHRLIA
1						AETGYSQSTVVRAFREAVNKGI
1		l				LSVEIVIGDHRERRANLYRFTPS
		ŀ				F\LAFAQQAKNALIESKLKISSA
1		l			1	ATKVKAVLAKTLALFNFLSTPP
1						CQNDTPSPCQDDVAIKNKKSQ
		l			1	VKKTKRSVSGGAGTTSLKKLTS
						WIAKAKAKADNLRLSKKRTQK
						HEFKQKVEAAARKYAYLKNKR
		l				SPDIGGISNFDNLPHCMTVNEA
1		l				LNAVLAKNKDNEQWGAVAGA
		ŀ				YIADITDGEDRARHFGLMSACF
						GVGMVAGPVAGGLLGAISLHA
1						PFLAAAVLNGLNLLLGCFLMQ
1						ESHKGERRGELLKIVVLPGDHV
ļ		l			İ	GQEITAEAIKVLKAISDVRSNV
		ŀ				KFDFENHLIGGAAIDLQGGGGK
24092	54460	Α	24230	1	1230	PRCVSKSLMLHCTR
24093	54461	A	24231	1	799	MKPRTYTVSVTAVKDGVSGVC
		1				SFRCSDVSRVSSFOWVRGLADF
		l				RSEAADLHNSGAQLASPIGSRT
		l				GATSGAPCQSRTHAPALLSLW
						AVDGTGRHGAGGSTRRQGSGG
		l				TQEPMVAGKAQAWRAAGFSG
		ŀ				SGDGVGVSILVPWGGTCIGSFR
						OANSWISGGAFSDAGCDSDRL
		ŀ				GKDWNSLEGSEEERKMWESLE
ļ.						LPRDLKSSEDRKMWESLELPRD
						LLNGFDQNAD\TICTMKSRLRW
		l				SQMEMKNLLGTGVNVTLVML*
						QRDWRHLLLV
24094	54462	Α	24232	416	775	
24095	54463	Α	24233	548	882	PGGHQYPPCHLEEKETDLPLLY
						TAVEEKGLSYSIIPGSLVEKPLT
				l		STNSSGTPIQSNITSL*IPLAGIMS
		1		l		MAASQN*AI*WPLAPTMQSFTC
						TT*RLS*SHRRSWWHRGCT
24096	54464	Α	24234	144	570	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	11100	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			,	sequence		, , ,
24097	54465	A	24235	254	1117	LSRMKLRTLVVSVTVLKDVVS
						GVCSFRYSDVSGVSSFQWVRG
						LADFRSEAADLRGLKLOTFVVS
						VTPHKGSVDPKSDQQRDLLQR
						VKEONFRNVKGDPAALAHEDD
				1		VNGTESGAPRKHGDRELDDVV
			ł			ITNPHIEAILENKDWIEDASGLM
						SHCIAILKICHTLTEKLVAMTM
						GSGAKMKTSAIVSDMIVVAKW
						NSPRVDDVVKSMYPPLDPKLL
						DTGTTTLLLSV\SHLVLVTRNA
						CHL\TG\GLDWI\DQSLS\AAEEH
						LEVLREAAL\ASEPDKGLPGPEG
						FLQEQSAI
24098	54466	A	24236	472	844	RTDGVLVWMSFLFVSFPSNSQD
						PQLQVCWSLLEVHSRPCLPGYQ
						QWRLLCPHEITGESMSKLHIIQK
		1			1	YTSSCCLFKSS\SERGSSTPCPSW
						LGSGEGLSTLHATDFSFSAPMA
						QRGKSSLVSLE
24099	54467	A	24237	206	456	
24100	54468	Α	24238	50	392	LGLLPFFHRCPTQRGGI*RGSLA
						TAALLSCGGLCPVGS\SRHLCL\
						PVRGKRPTQASVMGDALPPTK
						LECPRSASGCCAGSDNFKPVDL
		l				SLLSSMGLGSTELDHLAPWLQP
24101	54469	A	24239	1	204	PFQG SFSVOFCSLVGDPLG*SSCDPLE
24101	34409	I <sup>A</sup>	24239	l'	204	EKRCSGFWNIQPFCAGFSPSLW
						DYLSLVFDVDDLQMRSLSGCPF
24102	54470	A	24240	1	375	MCNAQVLEYMGKSSSLLTSDL
24102	34470	ľ	24240	ľ	373	OLVRDALRSLRNSFSGHDPOHH
						TIDSLEQGISSLMERLHVMETQ
						KKQERKPPLVIPRQT/WVWSGP
						PANSNRPAAEGPDC*KEN*RDA
						LRSLRNSFSGHDPQHHTIDSLEQ
						GISSLMERLHVMETOKKOERKP
				l		PLVIPRQTGSGVDLQQTPTDLQ
				ĺ		LRDLTVRRKTNEOKGIASASTK
						RTSTPKPHL
24103	54471	Α	24241	1	674	MVVYACSPSYLEAEVGGSLSLE
						VEATNPKNSSKKFLDLISEFCK
						VSGYKISVHKSVALLYTKNDQ
						AATQIKNLIPFTTAAKQFKYFGI
						YLTKEEERLKPSRKKITKKHTK
						KRTASLILHAMICSQIPKQQQNE
			i			KYQVPQFDQSTIKNIESAKGLD
						VWDSWPLQNADGTPPLVIPRQ
				ŀ		T/WVWSGPPANSNRPAAEGPDC
						T/WVWSGPPANSNRPAAEGPDC QKEN*QTERNSINIKKKDIHTKT

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
24104	54472	Α	24242	1676	1768	
24105	54473	Α	24243	270	471	PSREYQDANPWLKWIKYSVRT
		l				LNKSDCYASGYK*ACLLCSVYS
	1					HGKAAGECTLSAGSKNGLAEVI
24106	54474	-	24244	23	200	KK AOMKAHGRSRSS*RC\DEEHWR
24100	34474	l <sup>A</sup>	24244	23	200	ACRAQEPPCSQSQRERSKRTCR
1		1	ļ.			QGTVASKVKCHSEAR
24107	54475	A	24245	3	270	ASTSHVPQPICGLDDLAIRHSAV
24107	34473	ľ	2-12-13	ľ	2.0	RLNRERVHSTAVLP*EYTKQRR
		1				QGHL*PDPYKKEGAATNEDAL
		1				LLRWERGRGARLHIPGCKAPQL
24108	54476	A	24246	135	275	
24109	54477	Α	24247	365	583	
24110	54478	A	24248	2906	3114	
24111	54479	Α	24249	108	405	NLHTPYRPQSLGQGEPLDDYV
					ŀ	NTQGAS/LFSVTKKELGAGSIEE
1			1		l	CAAKCEEDKEFTCRYFHCRCTF
		1	1	İ		LSIPQVTSSFLCFPLPLPLLKSSK
L		Ļ.				LLANRDKYRM
24112	54480	A	24250	1	386 808	MSYSOCVSDPKAHGILPGYHC
24113	54481	I <sup>A</sup>	24251	1	000	WLFRDQGLLKSQHVINPAKAST
		ľ				GFFLODTLMVLEIKTTIRNYYK
		İ				HLYAHELERLEEMDKFLDTYT
						LRRLSOEVIDSPNRPIMSSKIESV
1		l				KGYTHOOFCHKSTLNKGDRVI
						DNLTCPPYCCVWFPLAGTGPHI
						LYLSRLVSNLELFKRGKGRGEQ
ł		1				RKEEVTCGMLRKSLVTSDLPEP
		1				GGVTSEGCETAKVTACSSLWE
		1			1	LCHREVLTPYCPKDTCRKWLE
	1	1				TPVRRSCPVRRSRISTCLRKKFG
						HILAEQLSCA*RCL**FL/YGCFY
		ı				FQDHECVLKEESSTSFGRINHV
						LTLKEPLVPE
24114	54482	Α	24252	833	1245	ANMFPAFAAVAAGAATEAAA
				1	1	AAAAAPAPLAPARRAAAAARG
		l		ł		PPLPRLP/WSCAEVMYHPQPYG
						ASQYLPNPMAATTCPTAYY/RA GAPTWPAGESPDGQPAGGGGE
						GHREYRSGARRRPAGTRGAED
				1		ALGSAGASLSRKP
		_	L	L	L	ALGOAGASLOKKY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	i	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24115	54483	A	24253	145	766	TGCIREKFHLPEGSLLIVSSRKH
		ľ.				LORLAFOAKVSSMMLPLOGAO
				1		MLOMLEKSLR\KSLPRNPLKVY
1						GT\VFHINHG\NPFNLKAVV\DK
						W\PDFN\TVVV\CPQ\EQDMDRW
						TLDHYTIT\YOIYSKDPONCOEF
Į.		1				LGSPELINWKOHLOIOSSOPSLN
		l				EAIQNLAAIKSFKVKQ\TQRILY
						MGRLETAK\ELTPFLAGNQRILS
						P\SGWAKPKAM
24116	54484	A	24254	215	550	VGFRSCLVMSGRGKGGKGLGK
24110	34404	^	27237	213	330	GGAKRHRKVLRDNIOGITKPAI
					1	RRLARRGGVKRISGLIYEETRG
				i		VLKVFLENVIRDAVTYTEHAK
						RKTVTAMDVVYAL\KROGRTL
24117	54485	A	24255	3	292	TLLNLAAAHCSGPCLLWFOLSF
24117	34403	^	24233	ľ	272	RSPSTSAVGRCRRPSADFHPSRS
						SRVSAVLLIQ/QRCPLPILIQLKA
		l		1		CHCSCTAKWLGSSPKPETFFEII
						HPSPS
24118	54486	Α	24256	1	789	111 51 5
24119	54487	A	24257	3	256	TLLNLATALSSGPCLLRLELSFH
24119	34407	l^	24237	ľ	230	SPSTTAVCCRHRPAADFPPFGR
		l				VSSLLLIQ/QRCPLPLLIGLKACQ
1						CSCAAKCCVRPNRAEQ
24120	54488	A	24258	3	212	CSCHARGE VIG MOLEQ
2412I	54489	A	24259	2	491	
24122	54490	A	24260	20	272	
24123	54491	Α	24261	160	876	
24124	54492	Α	24262	623	796	GANSWNLTGTLAVKAGTVENP
		İ				I*VFATHVESAGPR*LDPRTGSL
		1				AVRTKGRHWDQGA
24125	54493	Α	24263	242	659	TSNSPPOTRPHVCHGPRQYLLQ
						LSSRDLEFLVQIAVRSCLCPGLR
						IRSPVFAVASTSTFGTLRGKVV
						KDFGFVFPPPWGQPCLCPVEER
l	ł					ALSCOGPVVDSRCPRCGHEGM
						AYHTQQMRSADEGQTVFYT\C
		1				TNCKFQEK
24126	54494	Α	24264	66	238	,
24127	54495	Α	24265	2	278	
24128	54496	Α	24266	168	565	TENFRALSTGEKGFGYKSSCFH
					1	RIIPGFMCQGGDFIRHNGTGGK
				1	1	SIYGEKFDDENFILKHTGPGML
					1	SMANAGPNTNGSQFLICTAKTE
			1		1	WLDGKLVVFGKVNGMNIVEA
1						MECFVSRNGKTGQKITIA\DCG
24129	54497	В	24267	20	88	
24130	54498	С	24268	183	254	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, v=possible nucleotide insertion)
24131	54499	A	24269	412	590	KSGLATQG/GRRRGWVFLAPQ KGGEGVETQREGVGVLAPSPE
						KRDLPLRVKDQGRRPCVV
24132	54500	A	24270	550	852	
24133	54501	A	24271	207	381	
24134	54502	A	24272	412	589	KSGLATQG/GRRRGWVFLAPQ KGGEGVETQREGVGVLAPSPE KRDLPLRVKDQGRRPCVV
24135	54503	A	24273	191	235	
24136	54504	A	24274	95	375	AEKIWEGVSQRALGQSSRGSGS GCQMNPLSGRPQVVCFFTAHM LTKKREEGASMLNMPCFQ*KE EMIQANLGPIADSSHCLGIEGPP WAPLTP
24137	54505	A	24275	144	293	WQVPLFWQPFPEPLELWPKAL* LTPSQIFSD*RLKTDAARSPRKP PAPR
24138	54506	С	24276	1	909	
24139	54507	A	24277		1744	MGCLGTORAPRAFLLPLPLHE SGGGGERGATAAGTEQGAC LAASLLRGLWDAGEVTFIRVR SRPVDPTLAPLHEHSEAQCVAS PSLTVTKFKSLLPKSLLAGSRE PSLTVTKFKSLLPKSLLAGSRE RSRALLFOQARPPREPPQVTQ RTQLQGQRILQWRPATEGPHSP RRGSDSAKGTHPPTTGTCPPRR RQGAGYSRYNSGESRRLGRQ QDLVARSDEQRDPAPATRRED KMIEELQPFGEHRPRRSPSGQDL VLALCKAAAGPGAPPAFRRED TREHGYAWKLGDTRNHRALK RKSQTWLRRFPAGCPDVCSAL ADLRAFMGLRGEEVCADWSM GGHGSGECRGTGGQRAVVFK SPVQEKGKEPLSLLQYFMVDEB ERFLRITISHEDNDNSNSSNHS SINETKETCFHEPKTPAPVTDW EGSLLVFNNCRDASLHISCEFK SWGRGKYPSTSPSPSPLASPTFL SWGRGKYPSTSPSPSPLASPTFL SWGRGKYPSTSPSPSPLASPTFL GQGGELVTSRNAV GPGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ
24140	54508	A	24278	2	120	LTWQPLPEPLELWPKAL QSSAAKPRTSGKGSVREPWAR
-7170	15 7500	1	1-72.0	Γ	I	VPGALGVAAR*AEQSDFQ

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
24141	54509	Α	24279	1135	1483	ILTMTGSLSCTPSNAVIEKV*VC CSSLSTG*SKVQVASNTSLCDG ED*GVVIQKKVKLWVFCIFYVL VQSYYS*R*SQIETVRNGCSHW GRFQDIALITGHCKIWTLIVFIQ YFNV
24142	54510	A	24280	1	860	ATTGGTNLRDDMRLDDTVHU VIATPGRLDLIKKGVAKVDHV QMIVLDEADKLLSQDFVQIMED III.TLPKNRQILLYSATPLSVQK FMNSHLQNFVSINLMEELTLKG VTOYYAYVTERQKVHCLNTLF SRLQINQSIFCNSSQRVELLAK KISQLGYSGPYIHAKMRQEHRN RVFHDFRNGLCRNLVCTDLFTR GIDIQAYNVINFDFPKLAETVL HRIGRSGRFGHLGLANLITYDD RFNLKSIEAQL*TEIKPIPSNIDK SLYVAEYHSEPVEDEKP
24143	54511	Α	24281	848	1218	PMQRKSMLKISVS/PVSVQKFM NSHLQKPYEINLMEELTLKGVT QYYACITELQKLAETYLHRIGR SGRFGHPGLAINLITYDDHFNL KSIEEQLGTEIKPIPSNIDKSLCG AEYHSEPVEDEKP
24144	54512	A	24282	241	477	
24145	54513	A	24283	38	208	MTAGAPPPPPPFET\PEGV*GNP TCSGG*EDASGSTRSQRHPVGL PVSCALRACHLK
24146	54514	A	24284	794	1455	AQQGGTSKKRRFPODGOQRAG LACSRPPOPGCGCFQAPAGERY PNIKPHPTSVAPLVLCICYLMP MLPWAWSYWRGARRPSRHPG AFPCSVAHQCPSAARSPTLRQE DPEKGSQASSFPTGSPOGMVD RCLKGPQLDVSRKQLSLPQTGL RPVLVNQLSKKPTGYPQTQIQW VLMRDTGKDTDTEEKEGHAK MEAEIGVMYPQEHLEPPESGRA KGEIPL
24147	54515	Α	24285	90	354	QSPGVHQLSRFVS*SYSERP*SL LVSTRYHTRALH**HYADSIQ* ARSSKHTGLIGCTVP*RGFPRTH SFYSRKREPRRQTSSFRGIL
24148	54516	Α	24286	236	520	TACFLTHVAIKWVMHSSIPSSN GGGIYMIGLKAQVSYMRKWLK SPWSPLLLSPSLHRWPHGKFPM IS*QRKRRLGPGSQMVLHDMQ APPKSGQL

SEO ID	ISEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
110.	sequence	liou	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1	acquence		0,5,5,6,21,	sequence	or prepare sequence	
	ŀ					
24149	54517	Α	24287	16	278	FAQAMKPHLVQQLQLESPLG*
		1		1		AYDNPLSFSKIHLSDLHKPVL*L
						GDSNDVLGTLESMSAQSQCPV
		1		1		VPEMSDHLPFPMHSYPGKRPEV
24150	54518	A	24288	74	168	LGKCLFLHSCP*APPEAICLQLP
						LLSSPNGP
24151	54519	С	24289	92	564	
24152	54520	Ä	24290	1	876	
24153	54521	A	24291	71	282	PLVTQVVFRLFPTNSSLDLLQ/S
						PSP/GSGPHSNWQPFGSLDKWA
1	ļ		1		Į.	GVKHPNEECAASKIQTGPPCVV
		1				PLPWL
24154	54522	A	24292	81	1054	PRLFSDCFPPTAAWTCCRAFSC
	i .	1				SGPHSNWQPFGSLDKWAGVKH
					1	PNEECAASKIQTGPPCVVPLPW
						L*EEPNAAIYPSPVSPVCFLLLA
	1					HWIOSA*CHOCNGPVWYLVEV
	1	1				KSNQVLSK*HYDTKPES*YTPY
	1	1				GOEWRKGHLPSOWPHTRYOE
1						MY*FAQAMKPHLV*QLQLQSP
		1				LG*AYNNPLSFSKIHLPSPHGSH
1		1	l			SEPCL*GPTRTLV*L*L*KQTGM
		ı				LAVAPEENQHYLAWQLPQGRP
						SLLPOAVOGLSPOTKVERLMA
						ASVGEGMLPILGMEKLF/PSGK
1	l					KD*SGFTSSVSPASSGSSHTSPF
		1	1			QVAGSHSFPINAFTLTVDTW
24155	54523	Α	24293	5	88	`
24156	54524	Α	24294	779	1041	PPLSSPSGP/DEQSD/LWWQQSIP
		1	i .			LHQETVKVLH*MGPVPCATQL
						DKIHOOGFSDPYRGAVLLASG
1		1				WCPKRPEIPEEGATTHLCCFPAS
24157	54525	A	24295	248	3714	
24158	54526	A	24296	4177	7804	TFQESRFEQGFQKAIFILELVDA
						EDPPDEEEEASSIFSSSFHFLFP
	1	1				SSSSCFSFSSSSSSLLLRSPGDKD
		1				MPTAGMPSLLQSSSESPQSYPE
		1				GEDSQSPLQIPQSSPESDDTLYP
		İ			1	LQSPQSRSEGEDSSDPLQRPPEG
1			1			KDSQSPLQIPQSSPEGDDTQSPL
1		1	1			QNSQSSPEGKDSLSPLEISQSPPE
1					1	GEDVQSPLQNPASSFFSSALLSI
			1			FQSSPESIQSPFEGFPQSVLQIPV
	I		1		1	SAASSSTLVSIFQSSPESTQSP
24159	54527	A	24297	145	219	5. 2. 2. 2. 3. 1 5. 5. 1 Q 55. 2. 2. 1 Q 51
24139	34321	1^	27271	1175	217	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	delction, \=possible nucleotide insertion)
				sequence		
24160	54528	A	24298	191	1075	OPKVSRHCOTSPGSYLYSLOSG
24100	34328	^	24230	1127	1075	RWTTGKPLQQTKSVYLRSFHH
	i .					HHFONHRHYLHHHHLYHHHH
						YHPHPSHHHLHCY\SOHHHHH
	İ					HHHHHHHEHHLCRHOKSTOVE
i			l			YSVFAERENDLGLNGTGDEAL
						HDNGTTERVSGPPAGYLYRAD
						YAIGLQPFMPRLGRRTPGAGGP
						HSRAPELAYLQKLITSPATLPFV
						YLETGGWSPRDAHRVDNFDM
1						GPRENMGSVWEKSFLFTRNPSK
	1				i	SSYIADKNLIYLYASECIGRQSQ
	1				i	KARASKSGVQAEHPYITGAIKD
		L				DTRAIVKTRDLANRLK
24161	54529	A	24299	155	381	LICTSFKILSRLFCPSSL*SSSSQG
	1	l				ACISKLLSVLTYAHLYFALEKE
-	ŀ					VLTFLSRAIVSSSTLLTHGVAHS
						LTLFH
24162	54530	A	24300	1052	1338	TGWDGPPANCSSPMEGDSNVH
	1			l		CSCVCCCSCFGLLL/CAIHCCCA
	1	l				PPGCSACGAATLLYSAPLGSSA
	1	i				YGIALLSPFHHVRMMLSTKQEV
		i				GLHQTLNLP
24163	54531	A	24301	I	723	MLNVQALVVMVMCFMWASL
						WAPGGHPSAGNISSWVSGLVSF
					ĺ	PEWSSWELVVSVVGWAGKLW
			ł			LHESLLTLNLSCGLLVVLVALV
						FDEWHPWSGHVAEACSTGYWS
		l				TGGIWFHEIAAP*DCSSSPAME
	İ	1				QSWMEND\DELIEVGFRRLVIT
		1		l		NFSELKEDVRTHLKEAKNLEKR
	1	1		l		LDEWLTRINSIEKTLNDLMELK
	1	1		l		TMARELRDACTSFSSQFDQVEE
1	1	1	l	I		RVSVIEDOMNEMKREEKFREK
1	1	1		I		RVKRNEQSLQEI
24164	54532	c	24302	295	430	
27104	107002	<u> </u>	24202	270	730	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	coden for last amine acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
,				sequence		
24165	54533	A	24303	811	1947	OTERNSININRNGHOHONPICRS
2.1103	15.555	ľ.	2.000	ľ		PSSKTKGSIKPORSGRNOSRKA
				1		ENSKNOSTSSPPKDCSSSLAMK
			1	1		OSWMENDFDELTGAGFRRLVI
l				1		TDFSELKEDVOTHIKEAKNLE
1				1		KKIRQM\VTRINSVEKSLKDLM
	i			1		ELKTMARELCDACTSFSSRFDQ
		1		1		VEERVSVIEDOMNEMKRKEKF
		1		1		REKRVKRNEOSLOVIWDYVKR
ŀ				1		P\NLRLIGIPESDEENGTKLENTL
						ODIIQENFPNIARQAN/VOIQEIQ
				1		RTPQRYSSRRATPRHIIVRFTKV
		1		i		EMKEK\MLRAARQKGRVTHKG
	İ		1	l		KPIRLTADLSAETVQARREWGP
	1					IFNILKGKNFQPRISYPAKLSFIS
		1				EGEIKYFTDKOMLKDFVTTRPA
				1		LQELLKEALNMERNNWYQPLQ
24166	54534	Α	24304	<del> </del>	2394	EQUEENTER WILLIAM TO FE
24167	54535	A	24305	1902	2413	
24168	54536	A	24306	2	610	WAPPAIHIRCMPEAYNFPRKPL
		1				SLIPSAIQSNELIIMNGNRALNIQ
						LLKRYOCTLDIRYKOTSMLNC
				i		YEGTVITPSHIOYPCLSDSYGLY
						DIGIRNTDFLKGELSMNIRPLHD
						RVIVKRKEVETKSAGGIVLTGS
ŀ						AAAKSTRGEVLAVGNGRILEN
ŀ					,	GEVKPLDVKVGDIVIFNDGY\G
				1		VKSEKIDYEEVLIMSESDILAIV
24169	54537	A	24307	I	801	
24170	54538	Α	24308	1268	1579	TAIWSPSKSALYAAQTSGCSWI
				i		ALPSISTGSNAWIPRRCRVGARF
			1	1		SSTGCSRITSSRISQTTASSRSTIF
	i	İ		1		FAALMVVARPRSSSLP*MNGLN
1				1		SSSAIFFGRPH
24171	54539	Α	24309	583	796	IHICDIRFSATPAGGEFWQYIAS
		1			1	RQSIFHAIVETGHALIQPSSVPG
		1				LTPAMKAVATTSH*TVESCFRS
		1		ŀ		ASR
24172	54540	Α	24310	I	313	MPNVDVVGEMVNTMSASRSY
		ı		ľ		QANVEVLNTEKAMSIAVTTTDP
l						TNTGVSTTSSSSLTG\SNAADLQ
l		1				SSFLTLLVAQLKNQDPTNPMEN
			1	ļ.		NELTSQLAQISTTLRAY
24173	54541	Α	24311	54	323	SASAQVSALSIHSAAFQNEYLSP
1		1				CPN*RLV\SAFTNSSRQSG*PEK
1	1	1		l	1	SVSQTPVITASAFT*YADIAASV
						RNKILRPGTNVFGDTIGFRVQ
24174	54542	A	24312	816	866	`-
24175	54543	В	24313	1	985	
24176	54544	A	24314	1828	2505	
24177	54545	Α	24315	1	1938	

PCT/US01/08631

SEQ ID NO:	of peptide	hod	SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24178	54546	A	24316	4201	4668	
24179	54547	A	24317	269	503	
24180	54548	В	24318	1	1702	
24181	54549	Α	24319	829	1584	
24182	54550	Α	24320	1238	1926	IQAAYNRIAPSSSPKFPRACGK
	1			i		CLGERGRIRGPGPSFGLINRPST
	1					PDPTITGVSTT/SSSSLTGSNAA
	1	1				VQS/SFLTLLVAQLKNQDPTNF
				İ		MENNELTSQLAQISTVS/GIEKI
	I	1				NTTLGSISGQI/DNSQSLQASNI
		1			1	GHGVM/IPGTTVLAGTGSEEG/
						V/TTTTPFGVE\WDGTL/TDGST
		1		İ		APNGSYNVAIS/ASNGGTQLVA
		1	1			QPLQFALAAERFCALSPGAPYI
		1				YDTWVRQWVASP
24183	5455I	Α	2432I	487	1509	FYLQGLRQQPQVRQPLNIGVT.
		l				QGVNAAACHPHIAEQQLDHRI
		1		1		RTNVLRTNGVLRPAEGIQERG
		1				FIVSTGFCDVFADFQERVFWR
		1				TDVFNHIWRVAGNVLFQQVPI
		1				AARMLQRGIAFGEAVFVQFISI
		1				GGFIVLAFFRVVAGEQAIFKAV
						LTHDQAGGGIGFGVFAVEFFIV
			ł			QQVQQYA*QESNVSTGTNRGI
			l			VSYRRRTRKARIDHNQLRAVV
		1		İ		NFGFHRPAEPDRVSAHHHDEV
		1	ļ.			GVLDVDPVVGHRKTFFDIWHI
		1	ļ			DQFIDQRVRRFSGDNRRLQDR
			ŀ			QVGLQSANNSPIDKSLFRGINQ
	1	1	ŀ			QLYAAFTAQDVNIDIFKTRQQI
				_		FAVIGQTTGVQYGK
24184	54552	Α	24322	377	749	
24185	54553	A	24323	140	298	
24186	54554	Α	24324	342	1094	MGAPWLHRPRRALQRWRLGS
		1	l			GFSCGSGLGLQENNGGRPVMV
		ı	l			LALSTLPRRPTRCGCEFYCLIK
		1	l			GFRLGMKSVKMEIRGLWAGG
			į.			DKIAVPAEPRSAGPERTLSDLH
		ł	1			NESIF/ITGGGSGLGLALVERF/I
		ı	İ			EGAQVATLELSAAKV/ASLR/Q
		1				RFGEHILAVEGN/VTCYADYQI
	1	1	1			AVDQILT/RSGK/LVNTPAETLE
	1	1	1		1	TGFHELFNVNVLGYLLGAKAC
		1				APALIASEGSMIFTLSNAAWYF
5 1 1 OF		1.	24225	ļ	1505	GGGGPLYTASKHAATGR
24187	54555	Α	24325	I	1785	
24188	54556	A	24326	2	1760	TROUGO DE CONTROL DE C
24189	54557	Α	24327	2437	2679	TFQYIGQPHGDEFPHVAPVKH
		1	l		1	DKLGEGKNHHDKSRHLFHRFN
		1	1		1	H*MR*IRLTSNCAGSDAISQRD
	1	1	1	1	1	NKEKKNDQWHDTFNKG

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SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hed	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ł				sequence		
24190	54558	A	24328	3	415	
24191	54559	A	24329	ī	1932	
24191	54560	A	24329	1932	2117	RAISLGRTAAGYLNSDV AQRA
24192	34360	^	24330	1932	2117	ASGADPAGDFVVWH**IRKNLS
1				l		
		١.			1017	GGARYAIPHLYQHLAWHP
24193	5456I	Α	2433 I	1	1017	MRYREPPARGVEPNELHACRSE
						DKLSEDAFDDQCTGANPRYPLI
				l		SELKQILLDTYYGRDYVEDRLS
						FLLAVATLNGERLDGEMSEGK
			į.			LVMPFRHVSDAFEQTRETIGVR
						GNNAINDMVRQRLLN\RFTSEQ
						AEGNAIYRLTPLGIGITDYYIRQ
į				1		REFSTLRLSMQLSIVAGELKRA
	1	ļ	ľ			ADAAEEGGDEFHWHRNVYAPL
						KYSVAEIFDSIDLTQRLMDEQQ
		1				QQVKDDIAQLLNKDWRAAISS
	i	1		l		CELLLSETSGTLRELQDTLEAA
		1				GDKLQANLLRIQDATMTHDDL
	i	1				HFVDRLVFDLQSKLDRIISWGQ
1		1				QSIDLWIGYDRHVHKFIRTGID
				1		MDKKPRPSRRCL
24194	54562	Α	24332	5	220	GLSWLNYC*ASTLTISLRCATR
						AVPLTRIRCRPRLLPSRRERTAL
		1		1		PCIYVKIAVTLLTATCASCVRR
		1				WIPA
24195	54563	Α	24333	3	180	
24196	54564	A	24334	229	564	PGFKPRHSSWQRLPSLGHLPLY
						GSSVFTLLNLATAHSSGPCLLQ
						LELSFHSPSTTAVCRRHRPTADF
						HPSESGRVSHVLLIQ/QRCPLLL
						WIRLKACHCSCTAKCPGSSQLS
24197	54565	Α	24335	148	404	
24198	54566	Α	24336	213	1306	SCEHSVAHGRGIMDLSLLKALS
		1				EADAIASSEQEVRQIL\LEEADR
		1				LQKEVRVDGLGSVLIRLNESTG
						PKVMICAHMDEVGFMVRSISRE
		1		1		GAIDVLPVGNVRMAARQLQPV
						RITTREECKIPGLLDGDRQGND
						VSAMRVDIGARSYDEVMQAGI
		1				RPGDRVTFDTTFQVLPHQRVM
		1		1		GKAFDDRLGCYLLVTQLRELH
		1				DAELPAEVWLVASSSEEVGLR
	1	1	1	1		GGQTS\TRAVSPDVAIVLDTAC
		1		1	1	WAKNFDYGAANHROIGNGPM
		1	1			LVLSDKSLIAPPKLTAWVETVA
		1				AEIGVPLQADMFSNGGTDGGA
		1		1		VHLTGTGVPTVVMGPATRHGH
			1			CAASIADCRDILHMQQLLSAFI
		1	1		1	HKSNELGGVLRLRIQICRDGNE
24199	54567	В	24337	I	1075	THE SECOND PROPERTY OF THE PRO
24127	124301	10	~~1331	1	1.0.0	

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
24200	54568	A	24338	438	1462	QIVQCLDVPSYTFIWFAVLDPIY GDSYSTGLVYAIISIIVNAITIPIG LYLLNPSSGADGKKNSNLSALI SAAKEPVVWAPVLATILVLVG VIPAAWDPTINLIAKANSGVA VFAAGLTLAAHKEEFSAEIAYN TELKLILMPLALLLVGMACHLN SEHLOMITLASALPPAFSLKLIY CAAVARLMCYTRTGTTSLAITLE ETYTOKLTGSRPBISSYGRILM WPLAWYKDNGKPHLCSQLETT RHISTAAQGRSSGSLRRGLGAS PFYRGHDQARSEGQIPITIRIPMG LAQRAPKGRAOPRGLTAPFTN GALGGPQGRGTISKQASQTRPR ASAWPHITPI
24201	54569	A	24339	30	177	GSGYQRCHNNAASDRLRSYRK QPAER/WLAPPIDSITSPPASRRA LILLY
24202	54570	Α	24340	3350	3430	
24203	54571	A	24341	425	824	SISGSENACFNATNHSGRYSQY GACLTSSSCRKGQLARRGLSYQ SPHAPADPAFPAARRSPVFCLS DDGGHKSPRDMSPDKPSRHPD QVALTRRIPSPPAVALPSVVKR TVHEGGGDDDESNPSPLSPDVH
24204	54572	В	24342	I	2118	
24205	54573	Α	24343	1	888	
24206	54574	Α	24344	730	1101	
24207	54575	В	24345	1	1992	
24208	54576	A	24346	880	1103	TMMRRLISRQPVLPLQKCFSVLI SNPWRVFGGTGGGNQER/CRY AEESVAETRLCP*CRLYAEYDP GSGCAGLSL
24209	54577	A	24347	1	952	MITLEKLPLAVAVAAGVMSAQ AMAVDEHGVARSGIGWYGSGG EQQCFQTTGAQSKYRLGNECE TYAELKLQQEVWKEGDKSFYF TINVAYSVAQQNDWFATDPAF REANVQGKNLIEWLPGSTIWA GKRFYQRHDVIMIDFYYWDIS GPGAGLENIDVGFGKLSLAATR SSEAGGSSSFASNHIYDYS/NEN AND/YFDVRLARWDSFGGPLTG EPPSALACTSGVSGGKGYPGPP EAQVTFATPKKPGKEACSSIQV LNRGYRPFTESYNTOVRKSARD YYPRRQRSISITNSPSRRWKGKL NSRRCSTGNRPLVGAVS

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24212	54670	<u> </u>	24248	000	1274	ASFATTRPVTKLAMRVAIFDSV
24210	54578	A	24348	805	1274	VGORYWGSAASPEHSRFAVLW
		1				
						ILASIFR/DRIASLRVQATVKS*K
						MS*AWYWQGCFRNWKSSPA*Y
						QFFSPARSASSHFAGEPVSLHSP
						IALGYRSSESSSCHTSTASLSVH
						RDYAHLHGLRFDHSSSRRRLCY
		ļ. —			490	C
24211	54579	Α	24349	275	696	
	54580	Α		I	735	MFTPDYRSVSRVLGSGPGYASR
24213	5458I	Α	24351	I	/35	
						MNQPDVMPDIPGKRFYQRHDV
İ		1		ľ		HMIDFYYWDISGPGAGLENIDV
		l				GFGKLSLAATRSSEAGGSSSFA
						SNNIYDYTNETANDVFDVRLA
		1				QMEINPGGTLELGVDYGRANL
		l				RDNYRLGKGLSQGSGVAFDNE
1		1				KFAYNINNNGHMLRILDHGAIS
i		l				MGDNWDMMYVGMYQDIHWD
						NDHG\TK\WWT\AGIRP\MFKWT
		1				PIMNTGMKF/GYE/NVESQGPG
		1				QNIRTIYLAKP*QPDVMPDIPGK
		l				RFYQRHDVHMIDFYYWDISGP
						GAGLENIDVGFGKLSLAATRSS
		l				EAGGSSSFASNNIYDYTNETAN
						DVFDVRLAQMEINPGGTLELG
	ŀ					VDYGRANLRDNYRLGKGLSQG
		1				SGVAFDNEKFAYNINNNGHML
		1				RILDHGAISMGDNWDMMYVG
ļ		ł				MYQDIHWDNDHGHQVGGPAG
		l		1		IRPDVQVDAHHEHRDEIRYEKL
		_				NPRARDKTFVQFTSQNPDRC
24214	54582	Α	24352	1	3414	MIVQPLAIIGSHFDSIAAGNLAR
		ŀ				PIAVYGRNEITAIFASLKTMQQ
						ALRGTVSDVRKGSQEMHIGIAE
1		1	ŀ			IVAGNNDLSSRTEQQAASLAQT
		1		1		AASMEQLTATVGQNADNARQ
1		ŀ				ASELAKNAATTAQAGGVQVST
1		1		l	1	MTHTMQEIATSSQKIGDIISVID
	l	1	l	l		GIAFQTNILALNAAVEAARAGE
		1	1	l		QGRGFAVVAGEVRNLASRSAQ
				i		AAKEIKGLIEESVNRVQQGSKL
1		l		1		VNNAAATMIDIVSSVTRVNDIM
	l				1	GEIASASEEQQRGIE
24215	54583	Α	24353	1	2493	

SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24216	54584	A	24354	1	1201	MQPALLIVLNKVRQEARRAR
						ACRIRRASGIVQFGKVRRQLAA
						VELRQRQTPERFVLLRGTRQQA
	i					TGQRIVKTEQRVIIITQRSFRRPG
						EGRRINDHFRFPAHLPSIRPIGK
						HQTPFGISGHHFNFFTVAIGDDI
						AQLDRIGANQSVLKGFNKFVV
				1		QYATDSMTSQGKGLSQGSGVA
				ŀ		FDNEKFAYNI/SFNGHMLRILDH
		l		l		G\INWDNDNGTKWWTVGIRPM
		l		1		YKWTPIMSTVMEIGYDNVESQ
						RTGDKNNQYKITLAQQWQAG
						DSIWSRPAIRVFATYAKWDEK
				1		WRYDYTGNADNNANFGKAVP
						ADFNGGSFG\RGDSDEWTFGAQ
		l				MEIWGLSRASGTLVTRAWRYK
		1	1	İ	1	TTPTRHCLLMNHYAGEAAMR
1					1	WKNYLKSTQKVQLQNFANWP
		1				KYSLSTLFHFPFSKSC
24217	54585	Α	24355	188	406	FIRSFDPSFVPSVLDRFFVLSTVF
ŀ		1				HSSSIFLVVIDRPSCVGEPSLF*V
		1				GGSCPTNSLYRCSGLAWYLMR
						SMD
24218	54586	Α	24356	1	1798	
24219	54587	В	24357	1465	7841	
24220	54588	Α	24358	174	3762	TLNDAMRREKCLARINRTAKS
						GVVMPKLVTWMNNQRVGELT
						KLANGAHTFKYAPEWLASRYA
	l					RPLSLSLPLQRGNITSDAVFNFF
			l			DNLLPDRTIVRDRIVKRYHAKS
						RQPFDLLSEIGRDSVGAVTLIPE
		1				DETQHHYESLTITDEGEFVVFV
1			İ			GPSGCGKSTLLRMIAGLETITTG
		1	i			DLFIGEKRMNDTPPAERGVGM
						VFQSYALYPHLSVAENMSFGH
						ATGLAQHKEVAEVLPSRRICWI
		_				ANRKRSPVVSVSVWRL
24221	54589	Α	24359	364	845	RPLLDHGSSPRLNFLRFWYVVH
		l				AHTLMRMTEGWAGFDILRPYR
		1				WRRVIAFVVAFYFRHLIALLVV
		1				QMHFKLVKGNVDVVFTAQLID
			i			STVQLIDSTTTVTQFTHIQTQLM
						IQRAGTKCRKEDFRHWFVHHP
		1				WVL/MSPPSVTLYALFPDRSHR
						PRRDGSPDA
24222	54590	Α	24360	781	1275	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24223	54591	A	24361	1	1185	MDKFDANRRKLLALGGVALG
24223	34391	^	24301	1	1103	AAILPTPAFATLSTPRPRILTLNN
				l		LHTGESIKAEFFDGRGYIQEELA
1						KLNHFFRDYRANKIKSIDPGLF
						DOLYRLOGLIGTRKPSCGYVT
				ļ		TAPAARSYRLTSKFIAVGOKAL
				l		SSLNIIHIAAPHLEALIIGESDTK
						NFASHDHAILIYKLEPTTGMLR
						TRAYIGOHMPLYCSAMGKIYM
				ŀ		AFGHPDYVKSYWESHQHEIQPL
1				İ		TRNTITELPAMFDELAHIRESGR
			ľ	ļ		TMDREENELGVSCIAVPVFDIH
		l				GRVPYAVSISLSTSRLKO/PROH
1						RFRLREROPAGGTPPAGTNSVT
						GGASLHHARCORRK*KPYRPER
						A/ESFLPARGKICWP*AGTLTVC
						IMASASRLISGCTPDSTCWSTA
				ļ		HAPICRRRGRAINRCHCSS
24224	54592	A	24362	I	801	
24225	54593	Α	24363	1523	1950	LLPA/AQKRRVDFLRATDFNAL
						MPGVVEIDGKNIYTQIIDLTTRE
1				1		AVVNRPEVHRRYIDIQFLAWGE
1						EKIGIAIDTGNNKVSESLLEQRN
				1	1	IIFYHDSEHESFIEMIPGSYAIFFP
1						QDVHRPGCIMQTASEIRKIVVK
						VALTALN
24226	54594	Α	24364	281	997	LTFARLKGITRKIGRLQSRRRKR
1				i	1	CKAMGVDVNRVYSLDELVRG
i				1		NDILFSATGVTGGELVNGIQQT
1					-	ANGVRTQTLLIGGADQTPMAT
						LTEDDVLEQLDAQDNLF\SFMK
						TAHTILLQGIRQFLPSLFVDNDE
	1					EIVEYAVKPLLAQSGQIDDIDV
						ALRLIYALGKMDKWLYADITH
1						FSQFWHYLNEQDETPGFADDM
1	1			1		TWDFISNVNSITRNAMLYDALK
-						AMKFADFSVWSEARFSGMVKT

NO:	SEQ ID NO: of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino neid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24227	54595	A	24365	263	1410	AMISRRRFLQATAATIATSSGFG WHHY/CLNSDAIALGIEGRPDLI LLGGDYVLFDMSLNSSAYSDV LSPLAECAPITACFGNIBDRPUG TEKNHLIGETLKSAVITGMLNQ ATVIAYANRPFELVGTGDLWS CQCKSASASETNLSRLVLAHNS CSKEVMRDEPCDLMLGSHTHR WQLSVPMAGEPFCHVADERVV LGLNAFCERHIYTIPRGGKRIPQ TFAADLPCKGITVNPVQSTITEE TFQTLLVSRALEKLGYTVNKPS EVDYNVGYTSLASGDATFTAV NWTPLHDNNYEAAGGDKKFY DVHVGYTSLASGDATFTAV NWTPLHDNNYEAAGGDKKFY QYKITNIRQLEDPEIAKLVDTNG DGK ADLTICNPGLOFFGAINH QUCAYELTNTVTHN
24228	54596	В	24366	1	2872	
24229	54597	В	24367	1	1704	
24230	54598	В	24368	1	1470	
24231	54599	В	24369	1	2301	
24232	54600	В	24370	1	900	
24233	54601	В	24371	26	856	
24234	54602	В	24372	1	1743	
24235	54603	В	24373	1	1410	
24236	54604	В	24374	1	957	
24237	54605	В	24375	1	2742	
24238	54606	В	24376	755	1816	
24239	54607	В	24377	1	1968	
24240	54608	В	24378	926	6441	



SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24241	54609	A	24379	1	1739	MTWTHEAVNRREPLRGCEPNW
						DVDVKPAGCMISRRNELPYTLG
						LRIDMLVDLGTPASLMGQSSVY
						NTTDRHSTHRMDGLEQLMQQA
						DFAADIAVVAPWDNWSPTAGR
						FGLIRLLDEQTAPYAWYTSPMH
						LMDEKCSCHRQRPVIGLNQPQ
l			ŀ			YQSGTIEVDNVSFAYRDDYLVL
			l			KNINLSVPSRNFVALVGHTGSG
	ŀ		ł			KSTLASLLMGYYPLTEGEIRLD
						GRPLSSLSHSALRQGVAMVQQ
						DPVVLADTFLANVTLGRDISEE
						RVWQALETVQLAELARSMSDG
			İ			IYTPLGEQGNNLSVGQKQLLAL
						ARVLVETPQILILDEATASIDSG
						TEQAIQHALRRDCSPLIDHCDA
ľ						DTILVLHRGQAVEQGTHQQLL
					l	AAQGRYWQMYQLQLAGEELA
		ł		i	1	ASVQPLPNSDRRGSMKLVTVII
						KPFKLEDVREALSSIGIQGLTVT
l						EVKGFGRQKGHAELYRGAEYS
						VNFLPKAAYTGKIGDG/KIFVAE
				l	l	LQRVIRIRTGEADEAALYLWHT
l						ATERDGPFRNFTAGAFRASLPH
			l		1	PARHGRRPILAAPFMPVNPSRLP
					1	PRRRHLASGRSLRFNCCTAGDE
						TCSARSGERRSLVF
24242	54610	Α	24380	547	684	

	SEQ ID NO:					Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	lucation of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24243	54611	A	24381	l	2208	MLLVSASVNAMSISGQAGKEY
24243	34011	<u>۱</u> ^	24301	ľ	2206	TNIGVGFGTETTGLALSGNWTH
						NDDDGDVAGVGLGLNLPLGPL
						MATVGGKGVYTNPNYGDEGY
						AAAVGGGLQWKIGNSFRLFGE
						YYYSPDSLSSGIOSYEEANAGA
					ŀ	RYTICVQVFVVVMLIRNQLFTP
						VNTAIDGFATNYAIMPAYAVG
						EGVATHGLQFELMRTTGGNTQ
			1			HVFHHPLPLFAMWRYAAEMF
						YQQMCQFVRYHFFNKSFTVFQ
		l		l		QQNGIEANFIAFQPGRSGGSA
						LGRNVFGFRYQPQCQQDKQYV
		l				QSRIHPEGVGVTQRVEHGQERG
		[				ANDHVQSSGTELMQEGVMSAS
		1				LKNQQGFSLPEVMLAMVLMV
		l				MIVTALSGFORTLMNSLASRNO
						YQQLWRHGWQQTQLRAISPPA
į		l			1	NWQSGAAMLRVYHSNRLDVL
		l				EALMEFIVERERLDDPFEPEMIL
		l				VQSTGMAQWLQMTLSQKFGIA
		l				ANIDFPLPASFIWDMFVRVLPEI
		l				PKESAFNKQSMSWKLMTLLPQ
		l				LLEREDFTLLRHYLTDDSDKRK
		l				LFQLSSKAADLFDQYLVYRPD
		1				WLAOWETGHLVEGLGEAOAW
ŀ		1			1	QAPLWKALVEYTHQLGQPRW
						HRANLYQRFIETLESATTCPPG
						YLRASLYAAIQHALASVREHTT
		1			l	LVVIAHRLSTIVDADTILVLHRO
		1				QAVEQGTHQQLLAAQGRYWQ
					i	MYQLQLAGEELAASKGHAELY
24244	54612	A	24382	2	729	DCSPLIDHCDADTILVLHRGQA
24244	34012	^	24302	-	1/29	VEOGTHOOLLAAOGRYWOMY
l						OLOLAGEELAASVOPLPNSDRR
			l			GSMKLVTVIIKPFKLEDVREAL
						SSIGIQGLTVTEVKGFGRQKGH
ŀ		1				AELYRGAEYSVNFLPKAAYTG
		1				KIGDG/KIFVAELQRVIRIRTGEA
		1			ŀ	DEAALYLWHTATERDGPFRNF
						TAGAFRASLPHPARHGRRPILA
		l				APFMPVNPSRLPPRRRHLASGR
				I		
24245	51612	<del> </del> -	24202	683	1161	SLRFNCCTAGDETCSARSGERR
24245	54613	Α	24383	063	1101	RTGHCH/SAFSFSTVLTSCSLSRI
		1	į.			SFLAHRLQYLLRVAEPLPNSDR
				1		RGSMKLVTVIIKPFKLEDVREA
				1		LSSIGIQGLTVTEVKGFGRQKG
						HAELYRGAEYSVNFLPKVKIDV
	l .	ĺ		1	I	AIADDQLDEVIDIVSKAAYTGK
	I .	t				GDGKIFVAELQRVIRIRTGEADI

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24246	54614	A	24384	631	1149	FFGIKEOVITANNANSNRNKTD
						RCCRNWLEDHADNHCHKHSEV
						MPCLRRHTVGDRDKKHDTHH
		1			ŀ	HOWGO*FPYIIFWLVSPLIAISRS
		1				KVSTHGLRQRDFILLREASQSF
		l				RFVLRVYDVADHIAGKFTVHD
					1	LQFIGYHGGQAQLVAQVFSKES
	1	1			l	KAAGGNRHFPAORFOLEHOLR
24247	54615	Α	24385	833	1264	I I I I I I I I I I I I I I I I I I I
24248	54616	Ā	24386	1	1632	
24249	54617	Ā	24387	2078	2371	
24250	54618	A	24388	1	159	
24251	54619	Ā	24389	530	811	RPGTGTAAOVLNRRIKISRST*L
	0.015	ľ.			1	TVVMGYWSHPSVSLLKWDRR
	1	l		į .		KSPKCSSALRVNISNTLLSTARR
		l			i	RKRYLIVIPALLVVRRCRPRGPS
	l	1				ACG
24252	54620	A	24390	1806	2115	AAFOEIELAFODPOCVGLLTAP
27232	34020	ľ` <b>`</b>	2.550	1000		MLMPLFLDQLILQCFFW/W/TN
		1				LERA*TSFSALFIDLOPPGYRTT
	l	1			1	TSKHKVSSSLIKGHVLLDHSFH
		l			!	DLNTOLWISTNAFREGN
24253	54621	A	24391	1476	1751	ALLLVSFASORFISA/TVGNONA
24233	34021	l^	2-371	1770	1731	GVDLLNLTLLVGGIFVLNNAGN
						FAVFTGDTSITGRIIQFHRQQTN
	ł	l				ATLRFSITOTLECFDRDORHVA
	1		i			VEH
24254	54622	A	24392	606	4519	EEVVRAISAKRNEPEWMLEFRL
24254	34022	ľ	24372	000	4515	NAYRAWLEMEEPHWLKAHYD
		l				KLNYODYSYYSAPSCGNCDDT
	[	1				CASEPGAVOOTGANAFLSKEV
	l .	l				EAAFEQLGVPVREGKEVAVDAI
		1				FDSVSVATTYREKLAEQGIIFCS
		1				FGEAIHDHPELVRKYLGTVVPG
	1	l	ĺ	1		NDNFFAALNAAVASDGTFIYVP
		l				KGVRCPMELSTYFRINAEKTGO
	1	l				FERTILVADEDSYAAKRLTDYP
		ĺ		Į.		HQLSGGERQRVMIAMALLTRP
		l				
21255	54622	١	0.4202	120	256	ELLIADEPTTALDVSVQ
24255	54623	A	24393	129	256	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
24256	54624	A	24394	1	1082	MAMSKVKSITRESWILSTFPEW
24230	34024	^	24334	1,	1002	GSWLNEEIEQEQVAPGTFAMW
		1		l		WLGCTGIWLKSEGGTNVCVDF
	İ	1		1		WCGTGKQSHGNPLMKQGHQM
		1		l		QRMAGVKKLQPNLRTTPFVLD
		1				PFAIRQIDAVLATHDHNDHIDV
						NVAAAVMQNCADDVPFIGPKT
		1				CVDSWIGWGVPKERCIVVKPG
		ı				DVVQVTALEIHAPDAFDRTAPI
		l				TWLADOKRRRTVKGRTTRPCN
l		l				RSVTVRPRRRDAVIGRFYRHQS
1	1	l	1			ITQKGLLLMVMPGCAFPLWIDS
		l				ELNLTFIDALFEGVSGITTTGAT
		i .				VIDDVSSLPRAYLYYRSQLNFIG
	l	l				GLGVIVLAVAVLPLLGIGGAKL
		1				YQSEMP/AAI*G*QTHSPPGRYV
		1				TDTVDNLFFIRCRT
24257	54625	Α	24395	49	281	ASDHRYLCSOSVPLRGIPACYL
						LQAPLTRWR/RIPPSRRICAAIDS
		1				AISRSPQRWLTAVCCGTVNFTN
		1				TCLLPDGVCA
24258	54626	Α	24396	737	1419	CIWRMYSHRLVAWRRCVKRR/
		1				RYLTRFAKTRGVAIVMVGHVT
		1	ľ			KDGSLAGPKVLEHCIDCSVLLD
		1	i			GDADSRFRTLRSHKNRFGAVN
1					i	ELGVFAMTEQGLREVSNPRGR
					ł	GKRKFISPKKTPALAKFRRNLL
						PPKVPGNGTSVRFRHSRICWMN
						PLPGAAWNCILTLKKPISPGVQ
		l				VFPESFTPARLEAFAGRSQACL
1						VPAQNILHQKSQAWLFDFPEQE
						KELRELSAWMAGA
24259	54627	A	24397	526	1693	STTGIRSIAMSNPILSWRRVRA\
		1				LCVKETRQIVRDPSSWLIAVVIP
1		1				LLLLFIFGYGINLDSSKLRVGIFL
		1				EQRSEAALDF\THTMTGSPYIDA
						TISDNRQELIAKMQAGKIRGLV
	1			İ		VIPVDFAEQMERANATAPIQVI
		l				TDGSEPNTANFVQGYVEGIWQI
		l				WQMQRAEDNGQTFEPLIDVQT
						RYWFNPAAISQHFIIPGAVTIIM
		1				TVIGAILTSLVVAREWERGTME
1	l	1				ALLSTEITRT\ELLLCKLIP\YYF\
		1				LGMLAM\LLCM\LVSVFILGVP
	1	l		1		YRGSVLILFFISSLFLLSTLGMG
l	l	l				LLISTITRNQFNAAQVALNAAF
l		1				LPSIMLSGFIFQIDSMPAVIRAVT
1	l	1				YIIPARYFVSTLQSLFLAGNIPV
	l					VLVVNVLFLIASAVMFIGLTWL
	L	<u></u>		L	L	KTKRRLD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
ĺ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \≔possible nucleotide insertion)
				sequence		
24260	54628	A	24398	655	915	IAVNVTLAFNGGKOLR*STSLL
						CCQEGLAFAVNLRQHFRVQTIS
						FFFLPVLRFWRPGIQQAKKCHP
						APRTAKGIPQQVIRAMRLLFK
24261	54629	A	24399	549	827	
24262	54630	Α	24400	472	626	WRFFRPRKDNIRRTLFADGHQI
l						AAQPGLNLSCTRTIF*IKRINTFP
						GRWRK
24263	54631	Α	24401	333	617	SDQSETDYRFADGERHLENQPH
İ					\	AGQTGRGGAGPGCTLYQRFAQ
		1				LCEVSEVERSGCRTLSAQHVEQ
		1				NLPAAICGGDDADGAVVHLWT
		1		I		TA*RTRIGTLRSGPKMNDSAISII
		1				TATDSGWKILLHMLS
24264	54632	A	24402	74	265	
24265	54633	Α	24403	I	1005	
24266	54634	Α	24404	1068	1185	SVAIRQASTGNRCQCLCPACA
						WA*VNADVK*SARVSRL
24267	54635	Α	24405	1	2292	
24268	54636	A	24406	720	2496	RRAGYRKAADAMTTIVPEKLS
		1				REVGSHFHALDRTTAVHRQPVI
		1				SATTTVHEFLRCHQFPLVVCDG
		1		l		GRMTPRAMFLFSSAHILMPLNY
l					ļ	HNSHINTLFPVAGTLMVEPTES
		1		l		ESKVELDRFIDAMLAIRAEIDQ
l		l				VKAGVWPLEDNPLVNAPHIQS
1						ELVAEWAHPYSREVAVFPAAL
ŀ		1				VTGGSRGIGRATALLLAOEGYT
1				l		VAVNYOONLHAAQEVMNLITO
		ŀ			ŀ	AGGKAFVLQADISDENQVVAM
l		i i		1	ł	FTAIDQHDEPLAALVNNAGILF
		l				TOCTVENLTAERINRVLSTNVT
		ŀ			1	GYFLCCREAVKRMALKNGGSG
		l		l		GAIVNVSSVASRLGSPGEYVDY
		1				AASKGAIDTLTTGLSLEVAAOG
		l				IRVNCVRPGFIYTEMHASGGEA\
				l		GRVDRVKSNIPMQRGGQAEEF
				İ		AVOTFPVWRVAOHCAVRPFRO
			l	I		RIGOLRDIFHLKGDOFTNPRKT
						GVTPQPPCSTYPDPSPPPVPAAG
						HIDRPALAPTFTPPIMCSKATVK
						PLSPSPPGTPDPASVLPFREQRA
				İ		
l			1	l		AVHRLRTVVTRHLLALRRDPF
						ANRHRQQDQRNGGFHHRQRHL
l				1		HTGKTRSLHHHQFAALRQHPK
2 12 (0	51627		24407	40	106	TKQRRKQSRHREEDLHIFRHAQ
24269	54637	C	24407	49	186	
24270	54638	A	24408	470	811	
24271	54639	A	24409	1	1006	
24272	54640	В	24410	1	2793	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nueleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ı		sequence		
24273	54641	IA	24411	463	1258	CCRSSWCNSDSETGLQIVASYF
24213	34041	^	24411	403	1238	EDRNGDS\LALRPEGTAGCVRA
	ľ					GIEHGLLYNQEQRLWYIGPMFR
						HERPQKGRYRQFHQLGCEVFG
						LQGPDIDAELIMLTARW\WRAL
l .						GISEHVTLELNSIGSLEARANYR
i i						DALVAFLEQHKEKLDEDCKRR
	i	١.	1			MYTNPLRVLDSKNPEVQALLN
						DAPALGDYLDEESREHFAGLC
		l		l		KLLESAGIAYTVNQRLLNYRLS
						ANLIENNLSLFDGEGIYRFYHPK
				l		PTKNAERENRQCRNDLFFRCGF
24274	54642	Α	24412	22 I	832	TMAENQSTVENAKEKLDRWLK
						DGITTPGGKLPSERELGELLGIK
						RMTLRQALLNLEAESKIFRKDR
						KGWFVTHPRFNYSPELSASFOR
						AAIEQGREPF\WGFTEKNRTSDI
						PETLAPLIAVTPSTELYRITGWG
	i e				l	ALEGHKVFYHETYINPEVAPGF
						IEQLENHSFSAVWEKCYQKEPV
1		1				VKKLIFKPVRMPGDISKYLGGS
		1				AGMP
24275	54643	A	24413	158	481	7.0
24276	54644	A	24414	2201	2350	
24277	54645	Α	24415	62I	1830	AGWKRGGAGEPDQGLQA\EME
						VLENLQELKDLNQRQAHAESKI
						FRKDRKGWFVTQPRFNYSPELS
		ļ				ASFORAAIEOGREPSWGFTEKN
		1				RTSDIPETLAPLIAVTPSTELYRI
1	1	1				TGWGALEGHKVFYHETYINPE
	ì			į		VAPGFIEQLENHSFSAVWEKCY
		1			I	OKETVVKKLIFKPVRMPGDISK
		ı			l	YLGGSAGMPAILIEKHRADOOG
					I	NIVQIDIEYWRFEAVDLIINLCF
1					l	MVTINNARKILORVDTLPLYLH
ı		l				AYAFHLNMRLERVLPADLLDIA
						SENNLRGVKIHVLDGERFSLGN
1	1	1	1	1	I	MDDKELSAFGDKARRLNLDIHI
1	1	1		1	I	ETSASDKASIDEAVAIALKTGA
1	1	1			1	SSVRFYPRYESNLRDVLSIIAND
1		1		1	I	
1	1	1		i	I	IAYVRETYQDSRLTFNIEWGKS
1		1		I	1	SCCSPYVHPWAKLAAIYYRLQP HINDAF
24278	54646	A	24416	83	502	HINDAL
24279	54647	A	24417	15	95	
24280	54648	A	24418	500	786	LTORFGTVPLLLO\AYLYSAFFF
1	1	ľ.	1			TPEFLRPAPPLFSAAVLPEQRQR
1		1			i	AQGIGGLVIRPTVELHQALQLF
		1			1	FGHQPKIQSLHALLRGSTGFQA
	1	1			1	SFRLPL
		ь.		L		D1.40. D

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24281	54649	A	24419	1	2914	MYYSHASCVADQYACGMHRT
						PSRTGRPRIHAPRQARLKPQVK
						VASKKRNARDKMOPPNEOOKT
						RAHACTPSSRFRAVVRTLTCPV
						VANVTRIFPRYFSNTRVLRRRR
						KQARGTWWRLRHLHLYVQPH
						KHTSYYIYRLYRVTYVMSAQV
						VDTGIYEFTTRKHDIGTIELPNS
						EIRYHGNPAMSRPRKTPTPPNSP
1						ASERATGPTTLLPPIRORRYAA
						RHTSTIAYTNSTSDIFEAPYNKA
						APIMRYTSVYCSLFSP
24282	54650	A	24420	2	420	CPYPENRWRGKILARPLLFRAA
		ļ.,		ľ	1	*ASRFAGAGMPGRLIPTVRPRN
						SNSGRLVSASRCAKESLTVTTC
1						TPLPGSAVREASN
24283	54651	A	24421	212	497	TI DI GOTTI LETON
24284	54652	Α	24422	1	1149	
24285	54653	Α	24423	535	589	
24286	54654	Α	24424	1	1458	MFDPETLRTFIAVAETGSFSKA
						AERLCKTTATISYRIKLLEENTG
						VALFFRTTRSVTLTAVGEHLLS
		1				QARDWLSWLESMPERAATVW
		1				MFVPTPLSVASGLQRIGRAGHQ
		1				VGGVSKGLFFPRTRRDLVDSAV
		1				IVECMFAGRLENLTPPHNPLDV
		1				LAQQTVAAAAMDALQVDEWY
	1	1				SRVRRAAPWKDLPRRVFDATL
ŀ		1				DMLSGRYPSGDFSAFRPKLVW
		1				NRETGILTARPAAFAADSDIAD
i i	1	1			1	GQTQRFDFSILQSMAHDLSQTA
	1	1				WR/GIQYDAEKSLWHNVE\NRQ
		1				LDAQFFHMGMGFRRRVRMFSV
ŀ		1				DPATHLAREIHFRPELFKYNDA
		1				GVDTKQLEGQSDLGFAGFRVF
		1				KAPELARRDVVSFLGASYFRAV
		1				DDTYQYGLSARGLAIDTYTDSK
	1	1		1		EEFPDFTAFWFDTVKPGATTFT
				1		VYALLDSASITGAYKFTIHCEKS
1		1	ŀ			OVIMDVENHLYARKDIKQLGIA
		1				PMTSMFMPRKNASRLSKRCYDI
						RERDGREERK
24287	54655	Α	24425	157	650	PIRASTCLFKPLITQCFRQFGSE
1	1	1	l	1	1	GVNDHLLGILLIDPARAQIEYLF
1		1	l	1	1	IINTPHRRAVAAFHIICIDFQLRF
1	1	1	l	1	1	RIHFROFAHQQIVVGHLTIRFDG
1		1	l	1	1	VLRDVDQTVKYRAAFVADDSF
1	1	1	l	1	1	VQLSAVTKTFIVFQPGTGITHLV
		1	ĺ			FHRHRQAQ*PFKVPFSRRLRNGI
			l			нтнс
24288	54656	Α	24426	269	2024	
		-				

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24289	54657	Α	24427	1	3548	MPRSNYFSERCKGEIFLKFENM
		1				QRTGSFKIRGAFNKLSSLTDAE
	1					KRKGVVACSAGNHAQGVSLSC
						AMLGIDGKVVMPKGAPKSKVA
		1				ATCDYSAEVVLHGDNFNDTIA
		1				KVSEIVEMEARIFIPPYDDPKVI
						AGQGTIGLEIMEDLYDVDNVIV
						PIGGGGLIAGIAVAIKSINPTIRVI
						GVQSENVHGMAASFHSGEITTH
		l				RTTGTLADGCDVSRPGNLTYEI
		1			i .	VRELVDDIVLVSEDEIRNSMIAL
						IQRNKVVTEGA
24290	54658	В	24428	I	2691	
24291	54659	Α	24429	2174	3315	NLPKPDWARSAPHPASANYKS
						NTRRLPFAREQ/QDQHTLLAGT
l		1				LFIAKILIPMQIAQSAAYRRRCP
		1				DAQLHFLPLQLAPRGHRYKRV
l	1	1				CQVRYPPIIALIPVVFRLMNDAN
1		l				GLSQPADAGLAHHEMRNFLDF
1	İ					ASSNVAASRRRCVAALPCATG
1						KGRQANEQAENKCSKDPQTHR
						TDNKETKTPPARPTTTTQPHTK
		1				EEEPLSTPTQTKHPPTHVTGTGP
		1				HPTPASGQTEKHERKRPTQQKS
l						NARQNSKDKRTPNKSKTTKPN
	1					RGYNQHPAGEPTMSTHEKVKP
		1				HQQNRKIQAPPKRQKTDHRHT
						HNTHPHPPHRAASTTTQAAPPP
	l .					RRATRSGTSATAHNNTEPHGTR
		l		İ		HTQPGDHKRAEEHDRTADEPT
						PPEHTLPSHTQLGP
24292	54660	Α	24430	I	2925	
24293	54661	Α	24431	341	689	RPLAFRQRCTFYWR\PGSANCR
		ı		l		LV*GASGTDPRLYSPCAAAADD
		1			i	CGRRQNAGRRRRATSGD*SPHR
		1	1		i	RWENALLFDSRHRHCPRRAKN
		l				AGGEYRQRGIAGSDLQQRFTA
		ı				AEKDHSRS

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24294	54662	Α	24432	1	1931	MGTLLGTLTVLTLVAEELPAET
						DVAADPAASPTCTAVNQTIGRA
1			ļ		i	IVAADFFIIFQLWQNTVRQLVT
l						HFKPHGQSEMSDTPELINRARY
						LTYLYCRQPAKAIRELSEQELA
		l		l		WKKQATHEYQVLVTRCKQTL
		1				HQVEPLREIEPGRKRLVFDENL
						RPKQNPIPHLLDTCWPLTVDEH
İ						ANRSSIGNNNISFVVNRAVVTN
						VLNTMRTACNSQATIQIYNADI
						ARDFGTRGIFSINSGFSTVIRGG
			1			VVTVDLSAGAAIGDSALAFVID
1						DGLVAVNLDTDGLRIGDGHGA
						LVIHRCAVAGDINTVTAAADM
1	ľ	l	1			DLEDPRIISARFGVPRTQVRTW
			1			VALYEKHGEKGLIPKPKGVSAD
						PECVLGRESCDRAAHVPYQAA
					l	AHFMLAGSGSVARWLKVYEER
						GEAGLRALKIGTKRNIAISVDPE
				i		KAASALELSKDRRIEDLERQVR
						FLETRLMYLKKLKALAHPTKK
	1			1		AAEIPRSTFYYHLKALSKPDKY
				1		ADVKKRISEIYHENRARYGYRR
	1					VTLSLHREGKQINHKAVQRLM
				1		GPSHLKQRLRSSDTALTEERDF
		l	i	l	1	KATRPNEKWVTDVTEFAVNGR
		l	1	l		KLYLSPVIDLFNNEVISYSLSER
		l			1	PVMNMRRSRPGVGYFYALTPV
İ	1	l			I	YLANRLQALLGVEQ/HHPDGEL
	1			l	I	FLTGYVADGGYHS
24295	54663	Α	24433	3	158	
24296	54664	A	24434	1	3432	

3211

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24297	54665	A	24435	1298	2649	TFITRPTSNGTSIFKLMNRRRLE
	1005		2.133	1.20		GGNVATLR/RLTIGMIFQDHHL
					i	LMDRTVYDNVAIPLIIAGASGD
						DIRRRVSAALDKVGLLDKAKN
						FPIQLSGGEQQRVGIARAVVNK
						PAVLLADEPTGNLDDALSEGIL
						RLFEEFNRVGVTVLMATHDINL
						ISRRSYRMLTLSDGHLHGGFGG
						RLDRFRKSVGGSGDGGRNAPK
						RAKSSPKPVNRKTNVFNEQVR
						YAFHGALODLKSKPFATFLTV
		1			1	MVIAISLTLPSVCYMVYKNVNQ
					}	AATOYYPSPOITVYLOKTLDDD
						AAAGDPDRPSLQLRWRVIPGFS
						IAVAIYFLNESLGRPLMITGYSA
						ECSLLAWPVRSGLRKAPMKTFP
						LOSLTHEAQOKOFALVDSICRH
						FPGSEFLTGGDLGLTPATELKPR
					l	RLPSGVEQVLADAFHAQAAAL
					ĺ	VQGAGTGAIRAGLAALLKPGQ
					ĺ	RLLVHDAPVYPTTR
24298	54666		24436	546	707	KLLVHDAPVIPIIK
24298	54667	A A	24436	428	1464	RNSRSLAEISRLRGQTRSGLPM
24299	34007	A	24437	428	1404	LLNLOSMGAS/WYLSPVIDLFN
		1				NEVISYSLSERPVMNMDHHLL
	1	1				MDRTVYDNVAIPLIIAGASGDD
	ĺ	l				
		l				IRRRVSA ALDKVGLLDKAKNFP
		l				IQLSGACWHCRAVVNKPAVLL
		l				ADEPTGNLDDALSEGILRLFEEF NRVGVTVLMATHDINLISRRSY
		l				
		l				RMLTLSDGHLHGGFGGRLDRF
		l				RKSVGGSGDGGRNAPKRAKSS
		l				PKPVNRKTNVFNEQVRYAFHG
		l				ALQDLKSKPFATFLTVMVIAISL
		l				TLPTSCYMVYNIAKSERISDIST
		l	İ		ŀ	QLNAFPGCEVAVSDAPSGQLIV
		l				VVEAEDSETLIQTIESVRNVEGV
		ļ		ļ		LAVSLVYHQQEEQGEETP
24300	54668	A	24438	1	429	
24301	54669	A	24439	307	424	
24302	54670	٨	24440	300	468	V = GV-18 PPGP-V - 1 N 60
24303	54671	A	24441	946	1558	KGCVSIPFFSFSWA VMSNLTYL
		l	1			QGYPEQLLSQVRTLINEQRLGD
		l	1		I	VLAKRYPGTHDYATDKALWQ
			1			YTQDLKNQFLRNAPPINKVMY
		l			l	DNKIHVLKNALGLHTAVSRVQ
		l				GGKLKAKVE\IRVATVFRNAPE
		l	1			PFLRMIVVHELAHLKEKEHNK
	1					
	1	ı	l .			AFYQLCCHMEPQYHQLEFDTR
						AFYQLCCHMEPQYHQLEFDTR LCNRIRLNGHDHFRRRYALFRV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	eodon for last amino aeid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24304	54672	Α	24442	li -	774	
24305	54673	A	24443	1	452	MELKKLMEHISHPDYROAWKV
				[		EHKLSDILLLTICAVISGAEGWE
	1					DIEDFGETHPDFLKQYGDFENG
						IPVHDTIARVVSCICPAKFHESFI
						NWMLDYHSSDDKD/DHRN*W
		İ			l	KNTPAFL*QESP*GSDSCH*CVL
		l				NNAQSGHRTDQDG*EIQ
24306	54674	Α	24444	486	572	
24307	54675	Α	24445	1	683	MHSLVIGQIKTDKKSNEITAIPE
		l	ł			LLNMLDIKGKIIKTDAMGCQKD
	ļ	1				IAEKIQKQGGAVMNLQRFDDST
	1	l	Ì			LIRIFALHELHRLKEHGLTRGAL
	1		l			LDYHSRYKLVFL\AHPQPESRK
					i	LGPF\VADIHQWQNLDDYSPQY
		1				RQRVVVLLSHPANARDHTNVL
	l	1				MHVQGYFRPHIDSTERQQLRAL
		1				IDSYRRGEQPLLAPLMRIKHYM
	1	l				ALYPDAWLSGQRYFELWPRVI
						NLRHSGVL
24308	54676	A	24446	896	1066	WRLSPSSR*LMPKVPRIQAYFT
		l				T*YCS*VIQRSPC*FHCYSRSIPW
						AWGAGKQSLT
24309	54677	Α	24447	1	3660	
24310	54678	Α	24448	1	1530	
24311	54679	Α	24449	.1	1890	
24312	54680	Α	24450	746	1872	KHWRGSLISLSTPARFNGGPFW
		l				GANQGGDPKGRGGSQRALFRR
	i i			1		KNNRGGETVSGAEDACIVNNN
l				1	1	AAAVLLMLAATASGKEVVVSR
l	ŀ	l			1	GELVEIGGAFRIPDVMRQAGCT
l				ļ	1	LHEVGTTNRTHANDYRQAVNE
				1	1	NTALLMKVHTSNYSIQGFTKAI
				ł	1	DEAELVALGKELDVPVVTDLG
					1	SGSLVDLSQYGLPKEP\MPHELI
					Į.	AAGVSLVSFSGDKLLGGPQAGI
ŀ					l	IVGKKEMIARLQSHPLKRALRA
l	1	1			1	DKMTLAALEATLRLYLHPEALS
		l			1	EKLPTLRLLTRSAEVIQIQAQRL
		l			1	QDPLAAHYGAEFAVQVMPCLS
		l			1	QIGSGSLPVDRLPSAALTYTPH
l	1	1			1	DGRGSQLESLADRWRELPVPVI
L		ļ.—				GRIYDERLWLDLRCLEDEQRFL
24313	54681	Α	24451	1358	1419	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide detetion, \=possible nucleotide insertion)
24314	54682	A	24452		1568	MKGRNIDAQEIQKRYGMVGDP VIIGVVLQLIFGLAAGGEFKGC SALMITVAAIMVLFPRMIRLIVE GLLPISDGARKFFQKYFKGREV GLLPISDGARKFFQKYFKGREV GLPISDGARKFFQKYFKGREV FYGLDTAVTIGHPTTIAVGLLLI PIMLILASILPGNSVLPLADLPV APFFICMATVIHRGDLVRTLISG VIVMITVLLIATQFAPYFTEMAL KGGFSFAGESAQISALSVGNMF GWSISELMSIGIGRVVAVGIVA SVVLFLRVLSMSWLKEVIGTEK AVIAMCHLRALPGDPSFDAQLG MNWVIDKAWDDLMALQNGG VDAVMFSNEFSLPYLTKVRPET TAAMABIGQUMSDIRIPFGVNV LWDPVASFDLAMATGAKFIEI FTGAYASDFGVWDTNVGET HQHRIGAGEVKTLFNIVPEAAV YLGNRDICSIAKSTVFNNHPDA LCVSGLTAGTRTDSALLKRVKE TVPDTIVVLANTGVCLENVEEQ LISADGCVTATTFKKDGVFANF
24315	54683	В	24453	1	3352	VDQARSNNPSAIPAAFVKSIDG DRYQINPNRHGVNDG
24316	54684	В	24454	60	2284	
24317	54685	В	24455	1	2475	
24318	54686	В	24456	1	2890	
24319	54687	A	24457	1065	1139	
24320	54688	A	24458	282	615	
24321	54689	A	24459	2657	2919	
24322	54690	Α	24460	1	835	
24323	54691	Α	24461	167	338	
24324	54692	Α	24462	834	ISIS	LRWCSMPCCRWSKDRAAPSPA YSRLSSHVIIAGYVGGGVARG' HSWVFVALTLVLTAVLFSLAGL LNGVFAKTFDDISLVPTFVLTFL TVLGGVFYSLTLLPPFWQGLSH LNPIVYMISGFRYGFPWLNHNR PGVAVDINRLLLPRKQEKLGIK LDKDQLLAGVQDAFADKSKLS DQEIEGTLQAFEARVKSSAQAK MEKDAADNEAKGKEYREKFR QIPSVSSCOKQ
24325	54693	A	24463	3	557	
24326	54694	A	24464	255	693	
24327	54695	A	24465	3	242	QKAPVLANQRQPRRPTGALLLE /SLPLVAESSDG*EPMTNATEPH SRPLP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nuclcotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \*=possible nucleotide insertion)
24328	54696	A	24466	1	2636	MKNIIRTPETHPLTWRLRDDKQ PVWLDEYRSKNGYEĞARKALT GLSPDEIVNOVKDAĞLKĞRĞG AGFSTĞRK WILMPKDESMNIRY LLCNADEMEPĞTYXDRLLMEĞ LLCNADEMEPĞTYXDRLLMEĞ LPHLLVEĞMLİSAFALKAYRĞY IFLRĞEYIEAAVNLRRAİAETTE AĞLLĞKNIMĞTĞFDFELFVHTĞ AĞKİYCĞEĞTPİLLKPĞTĞĞWÜĞ RHSCWKLQELAMNDRDFMRY SRQİLLDDİALDĞQKİLDSÖV LIDİĞLĞÜĞDYLLDÖĞVLLDSÖV LIDİĞLĞÜĞTAPALYLA
24329	54697	Α	24467	506	673	
24330	54698	С	24468	1	3411	
24331	54699	Α	24469	2455	2694	
24332	54700	A	24470	93	1081	MGGIRDPWLHKLRSLWGICE YGHKEGYDHIRELNLAPGOTT LVPTGLAIHIADPSLAAMMLPR SGLGHKHGIVLGNIVGLIDSDY GQQLMISWWNRGQDSFTIQPGE RIAQMIFVPVVQAEFRILVEDFD ATDREGGGGISGRYFCNED KOTAKRNRREEILQSLALMLES SDGSQRITTAKLAASVGVSEAA LVRHFPSKTRMFDSLIEFIEST TRINLILKDEKDTTARLRLIVLL LGFGERNPGLTRILTCHALL EQDRLQGRINQLFERIEAQLRQ VLREKRMREGGGYTTDETLLA SQILAFCEGMLSFFVSEFKYRP TDDFDARWFLIAAISCSIMTPD TDGFCKADAATGKSSDMTD TDVYAGKSGRIPS
24333		A				RCKCADAATRGKSRRDLPDVQ NSARQYG/W/GCATHARSG+HI VEAVQSG/TPPQPLRPRGTSRKV STDRYWRAASQLPAQYQDLTSI YDVYTVPDRVAKRAFKSPPDD GPVMWRMGHQPHLIPRNVLAQ
24334	54702	A	24472	254	1150	RETICRYOPOMAEDMOSLVOG TYTRSG'LSVVNNNNGELLAK ELSVQVGIRPEYLDRLLHAFSG GQRQRIALARALSSQPDVIVLDE HGLTYVLISHNVSVIRHMSDRV AVMYLOQIVELGDAQQEIAQA ANLKLESKLAIMEQVYGKKVID AVIVGPKVDVSAVKERIVIQEV LEASDIPYRHDRQLLHNALEKA LQALGGLEVIQLYRDSNLVPG QLKTSEMGKTRLYAADSRLIA AKGSPIQPTLDSLK

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
24335	54703	A	24473	1874	2927	KEDOVKGGMVCPKKLLCOGTR
2-1555	34703	l'`			[-/-/	*RGGVVGGISQASVTSVIGYSA
		l			1	SCOPTPAIMOLHL/ANAASRRL
		1			ł	OPDRNRPNPGWHRPPIHAFTTP
		1				GLVSVSFLGYPCSTPTCLPFSES
		1			ł	SVGESVNLWPOSNENLRRTLV
		1				GEGNLLLTFIGNGKHEIIASIFFA
		1				FSEGISAQSHWQPRYISPASVCT
		ı				LVTEIDIKANKITFSIFDENAGA
		i				WLPPFCKAVVPGWVYKPNLGH
		l			1	EIACVDHWLMPNRLIENLCDIM
ł		l				LPIWVLTKFNDCCAGCVPCCVII
		l				VDDALHGAPKRMCCCITRGMR
		l			1	LLKLSVHAATRSTLQQASHGH
		l				CAWSAPGWLDTSGMDGDRLW
		1				WTLHEVLRDLCTSHGNIVESAR
24336	54704	В	24474	1	672	
24337	54705	В	24475	1	2496	
24338	54706	A	24476	3260	3363	RPDSGGP*RQRRYLRCADHRRR
		_				AVRNQQQLTACA
24339	54707	A	24477	112	224	
24340	54708	Α	24478	1614	2326	CTYSRKCRPDKRSASGNFALSS
1		l				LKSSPLTKNRRYWKVSPTHLLL
l		l				SKRPKISLNAYFGACSLVRPVL
		l				AAASDWTLQGAWSPTRAKALP
		l				ICESWRIPYADSLSSLAASCDAV
		l			İ	FVHSSTASHFDVVSTLLNAGVH
		l				ICVDKPLAENLRDAERLVEG\M
		l			l	REWREERGQGVVHKPIPGWQS
		l				TLEQRGFVGCARHFIECVQNQT
		l	İ			VPQTAGEQAVLAQRIVDKICSS
		_				FNVVRVAPLGDPIQSKPVV
24341	54709	A	24479	19	885	
24342	54710	В	24480	1	1704	
24343	54711	A	24481	974	1156	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Step codon, /=possible nucleotide
1	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24344	54712	Α	24482	162	1602	HRRKRARGTPKATTTRKPARTI
		1		1		ETLSPLLAILAFAPKLAICTRSPL
	1			1		L/DDMTPFTQCTNAYSHSGIRNP
		1		1		TAISSAAYPRRITGY/KLGPTGE
	İ	1				ERTVYGIYGISESNLAECEKGV
1						KSAVALTPALOPIDGVAVSYID
1		1				AAVALGNTINEMDKYYTQENY
		1				KDDAFAKGKTLHOTFLKNLEA
						FEPVAESYHAAIQEINDKRQLA
						ELKNIEEREGKTFHYYSLAVMI
1						SAKQINNLISQDKFDAEAAMKK
						VSELETLVAQAKEADKGGMNF
		1				SFINSAGQYQLEAKKYVRRIRD
		1				KVPYSDWDKEOLODANSSWM
	1					VEDSFPRALREYNEMLMTRIGG
1						GAGDODMVNDFSYSPOLVGRG
1	l					
		1				WFNDQVATGCTRDSTFNQQQV
		1		ì	ŀ	ALSVNAYYVVNRFGKIDWKPM
						YAEADKLLAKPNMRFKSDKLV
1	1	1				GDLSIGDQQMVEIAKVLSYESK
		1		i		VIIMDEPTDALTDTEAESLFRVI
		_				REMKSQAPRYCLYLPPHENNL
24345	54713	В	24483	1	1263	
24346	54714	В	24484	148	894	
24347	54715	В	24485	1	1071	
24348	54716	В	24486	168	1088	
24349	54717	В	24487	1	1236	
24350	54718	В	24488	1	849	
24351	54719	В	24489	1	1548	
24352	54720	В	24490	1	1372	
24353	54721	В	24491	1	2469	
24354	54722	В	24492	1	1917	
24355	54723	В	24493	1	1159	
24356	54724	В	24494	1	2294	
24357	54725	В	24495	217	2476	
24358	54726	Α	24496	477	916	SLPECEDRVQYRRSLFADPSYR
1		1		i		YGLARHGDPRRAGRLRDV*PL
		1				HE*YRHWHDQS/RLLIASVLPQ
1				l		VWHGACNAMTSSCIARQVS*N
		1		l		TALIIRMRKGDLSAVMSSQTGL
		l		I		INIVRRGLLLSCFRLTAMKISTV
		ı		1		SPFRPQMPSIVRSVWC
24359	54727	Α	24497	1136	1317	
24360	54728	Α	24498	3	317	
24361	54729	Α	24499	1	1245	
$\overline{}$		_				

SEO ID	SEO ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24362	54730	A	24500	85	1293	ITVATTQHQAGLTLRRAGIRQP
						DMAVORDHLPFRVAGVIRHIM
	-					LMLHSORRTGLLITAKNRLLFF
						ORRSVPKFHSKLNOAYCLIAAT
						DCPRRLLLICRYPHHFLHQLIIL
	ŀ		l			AIGIGSAVFEPHQIIGVSRCCPR
						HLGVERVLFLFLNDFIEOVIIRC
						HVVCQSAPAAANIQYAHPWF*
	ŀ					LOFFTNHFELRLLGRFOIIGIFPV
						SAGILHEGIKHAAEQVITQIVML
				i		FADYPGTFFTL/RD*RDERROY
				i		AACL*NDGGVGSRDLRVRLC*
	l					KTYPVGHIPTSHPYN/CSPNPKE
			Ì			PSCRTRLKKSGFFTRISAGAEPL
						MAIPARLSNSLACGARDSAAKE
						VLLWFRFVLLLKQTLLNWTTK
						NVTSLWRTLGSPVGVKEVRAW
				l		TIPV/VSNRAAGSGQNPY*F*KR
24363	5473 I	Α	24501	2794	353 I	SPRDSPTFWTKGGPGGSRGKAL
						RLVVGTDFHQAADNIPDHMMY
						KSIRFNIYHHIRTVTRDINMHNV
						APRGSGLALHRTERGEIILAQQ
		1				ALRSPMHPFSVQRFVKMGYPFT
		1				QYRRAQPMIVDDVAITASRGA
		1	ŀ	ł		KSGMKIIRYSFNPTHSDVTR*IRI
	i			1		YAQGPRRVFAQRGCFEMHNLP
						GTVYPCVSTPGAEDGDGFVRH
		1				LGERLFQLLLHTADFVLPLPAVI
						LTGGVLNAQRNFINRRLLPYVE
24364	54732	A	24502	1	1767	
24365	54733	A	24503	1	1275	SNPTAIGGRPRRPYQAEDRSLD
		1				FVDTALNRKILKDSIFHALQTIM
		1				VGIKDFLRLTQVFFDLATRIPRH
		1				LYHPVDITTHYGRFCRHRRHHF
						QLLQLCFRFLFCLFRHLRRVDF
			ŀ			ALQGFVFVRRVVHFTEFFLNSF
	1			l		HLLVQIVLTLGFLHLLFNAVAN
	İ		1			AFLNLQQIDFRFHHCHQIFQTFV
				İ		NVGHLQNRHRKIGIVIFRSDVL
				l		EHPHRNNTIKLVVQVTIILQQN
						GDIQAFTAFLSHFLLFGRNGDA
						HHAYVIIRCHVVCQSAPAAANI
						QYAHPWF*LQFFTNHFELRLLG
				1		RFQIIGIFPVSAGILHEGIKHAAE
			1			QVITQIVMLFADYPGTFFTLQIE
				I		ETSAGNTQRVFEMMGELVLET
						RLRSPAGFTQWIPHQGCRWSCL
		1		I		PVPRGTLALLSPWVVDGTGRR
24266	54734	١.	24504		2052	GAGGGTPQGGSSCTGAHGGGG
24366	34/34	A	24504	I	2032	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nuelcotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24367	54735	Α	24505	1	1060	MNARQQVAALRVPGVEKATIV
						YRRSLQEMPAWREEYEEALHD
		1		İ		GFRFLNNPERFDADGTLTLRVM
		l				SLGEPDEKGRRRPVETNETVTL
				l		LVDSLITAIGEQQDTEALNAMG
				l		VPLDKNGWPDVDHNGETRLTD
		l				VFMIGDVQRGPSSIVAAVGTAR
l		1	İ			RATDAILSRENIRSHQNDKYWN
		ł				NVNPAEIYQRKGDISITLVNSDD
1						RDAFVAQEAARCLECNYVCSK
						CV\DVCPNRANVSIAVPGFQNR
						FQTLH\LEA\YCNECGNCAQFCP
		1				WNGKPYKDKITVFSLAQDFDN
ļ	}					SSNPGFLVEDCRWVPHRGLYH
l						RLYREPTWRSEWPRRHICPEGS
	Į					TFPERHCDHLSANNSRNKLLPL
		_				QKPGFIN
24368	54736	Α.	24506	1316	2678	NUMBER OF THE PARTY OF THE PART
24369	54737	Α	24507	2172	5242	IYFSERLPALCTPDAHYREPCQ
l		1		l		YNCTRLDYDSALNIRELKKVAL
ŀ	ŀ	1				EKGWDEYKQRWHKPAGSGSR
	ļ	1				HPVAVIGAGPAGLAAGYFLAR AGHPVTLFEREANAGGVVKNI
l				1		
		l				VPQFRIPAELIQHDIDFVAAHGV KFEYGCSPDLTIEQLKNQGFHY
				1		VLIATGTDKNSGVKLAGDNON
		ı		Į.		VWKSLPFLREYNKGTALKLGK
1		1				HVVVVGAGNTAMDCARAALR
l	1	l				VPGVEKATIVYRRSLOEMPAW
	1	l				REEYEEALHDGVEFRFLNNPE
24370	54738	Ā	24508	192	484	RGLVHRGRYRWRQKWSAGSP
2.0.0	5.750	· ·		1		ARHPV/SQSTTDTPATHALSTKII
l		l				SPIPNAPISALARTTATRL*RSM
		1				NQPPVKRPTVIPRVKPA*ISEPE
		l				VGPIILGP
24371	54739	Α	24509	1	1713	
24372	54740	Α	24510	I	651	
24373	54741	Α	24511	3	783	PVLSKTPLTAKAIDAAQPQDKP
		l				YKLTDSLTPGLFLLVHPNGSKY
		1				WRFRYWLNKREFLQAIGVYPLI
ŀ		1				TLKEARRRATESRSLIANGINPV
						EQARKEKAIDALNMAAGFKKV
!			1	1		AEDWFATRVGGWSESYAKQV
1	1		l	1		RSALEKDVYPVLGKRSIVDITA
1	1	1	ŀ	1		RDVLALLQKKERTAPEQARNV\
1			l	1		AGASVRSSSLPLSPNWGGWLA
1						RAAFPHHITIAARMTKHPPCW
1			1	1		WCTILACRQASLAVRGSTHYSL
		L				ELLIRRHILSLLRSPICASPLTV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24374	54742	A	24512	746	2387	WTALPARLKQIRSGRKLPLTVM
		1				PALRETDTFDTFMESSWYYAR
		1		İ		YTCPQYKEGMLDSEAANYWLP
i		1				VDIYIGGIEHAIMHLLYFRFFHK
	1					LMRDASMTNPSSVPRAGGPAL
	1	1				LFENPKGYSMPVLCNLFGTPKR
					1	VAMGMGQEDVSALREVGKLL
		l				AFLKEPEPPKGFRDLFDKLPQF
	ı					KQVLNMPTKRLRGAPCQQKIV
						SGDDVDLNRIPIMTCWPEDAAP
	i			i		LITWGLTVTRGPHKERQNLGIY
				l		RQQLIGKNKLIMRWLSHRGG\A
				l		LDYQEWCAAHPGERFP\VSVAL
	1					GAVPATILGAVTPVP\DTLSEYA
						FAGLLRGTKTEVVKCISNDLEV
	1	l			1	PASAEIVLEGYIEQGETAPEGPY
	1	1			-	GDHTGYYNEVDSFPVFTVTHIT
						QREDAIYHSTYTGRPPDEPAVL
		1			l	GVALNEVFVPILQKQFPEIVDFY
		1				LPPEGCSYRLAVVTIKKQYAGH
		1				AKRVMMGVWSFLRQFMYTKF
	Į.	1				VIVCDDDVNARDWNDVIWAIT
		1			l	TRMDPARDTVLVENTPIDYLDF
	i					ASPVSGLGSKMGLDATNKWPG
						ETQREWGRPIKKDPDVVAHIDA
24375	54743	A	24513	576	878	NLGSVWHYFRRSPEPATHHDA
					1	ANLERSPAA*RLSGARYRILAV
	ŀ					*ADQSQTGSGDGSPDLQTGRV
					İ	GDEARHRKRGLHVPQPRSEPPV
						GARGFPYRFATRWR
24376	54744	Α	24514	2	878	
24377	54745	A	24515	1	735	
24378	54746	A	24516	I	1992	
24379	54747	A	24517	721	1239	

3220

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ŀ		1		sequence		
				l		
24380	54748	Α	24518	1	1954	RQLNGVQLAFRESFHTVTRVTE
		1				QHAAGAVAIHQHRNQLLTGAL
1		ı	1			GIFAVAVCCLQQRLDILLADHL
		l				AEHVQFVIAQFVTCQQHGNGV
		1				RNRTIIFLLFNELRKIVETVRVQ
						QTQTGEVAFHPQLFRGRVVYG
		1				PDLNARRGNGVINCSTLEDPQF
		1				GLILFVYTIGIQVGPGFFASLRV
		1				SGLRLNLFAVLIVIIGGLVTAIL
		1				HKLFDIPLPVVLGIFSGAVTNTP
1		l				ALGAGOQILRDLGTPMEMVDQ
						MGMSYAMAYPFGICGILFTMW
		1				MLRVIFRVNVETEAQQHESSRT
		1				NGGALIKTINIRVENPNLHDLAI
		1			ŀ	KDVPILNGDKIICSRLKREETLK
		1				VPSPDTIIQLGDLLHLDSRGSPL
		1				DTQDIDLQINSSVEKQFGDAIRT
		1				TILDVLARYNVRGVQLNVDDK
		1				GALDCILRARLEALLARASGLP
						MISASLOORKTRTRRSMLFVPG
						ANAAMVNNSFIYPADALMFDL
						EDSVALREKDTARRMVYHALO
		1				HPLYRDIETIVRVNALDSEWGV
						NDLEAVVRGGADVVRLTKTDT
		l		l		AODVLDIEK\EILRIEKACGREP
	İ	l			1	GSTGLLAAIESPLGITRAVEIAH
		l				ASERLIGIALGAEDYVRNLRTE
		l				RSPEGTELLSARCPIWQAARSA
						GIQAFDTVYSDANNEAGFL/QE
24381	54749	Α	24519	5121	5492	
24382	54750	A	24520	2520	3525	
24383	54751	A	24521	I	1263	MAINSPLNIAAQPGKTRLRKSL
		1		1		KLWOVVMMGLAYLTPMTVFD
		1				TFGIVSGISDGHVPASYLLALAG
		1			i	VLFTAISYGKLVRQFPEAGSAY
		1	İ			TYAQKSINPHVGFMVGWSSLL
						DCLLAVVIIAAIYVGFSLAGFHV
l						YIGMLEVPGANAAMVSNSFIYP
						ADALMFDLEDSVALREKDTAR
						RMVYHALOHPLYRDIETIVRVN
1		1				ALDSEWGVNDLEAVVRGGAD
1		1				VVRLPKTDTAQDVLDIEKEILRI
l		1				EKACGREPGSTGLLAAIESPLGI
						TRAVEIAHASERLIGIALGAEDY
1						VRNLRTERSPEGTELLFARCSIL
1					1	QAARSAGIQAFDTVYSDANNE
						AGFLOEAAHIKOLGFDGKSLIN
				1		
		1		1		PRQIDLLHNLYAPTQKEVDHAR
1					1	RVVEAAEAAAREG\LGVVSLN
2.120.4	5.4750	<u> </u>	24522	-	405	GKMVDGPVIDRARLVLSRAELS
24384	54752	В	24522	62	1403	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1		l		sequence		
24385	54753	A	24523	891	3240	RYRKSPGSWR/CMSCILTSWITT
24303	34733		2.4323	05.	52.10	MFPI*RLGQWC*NPCVTMAL/L
		l				APIDVHLMVKPVDRIVPDFAAA
ŀ				İ		GASIITFHPEASEHVDRTLOLIK
		l				ENGCKAGLVFNPATPLSYLDY
						VMDKLDVILLMSVNPGFGGQS
	1			l		FIPOTLDKLREVRRRIDESGFDI
	1			i		RLEVDGGVKVNNIGEIAAAGA
l		1	İ			DMFVAGSAIFDQPDYKKVIDE
		1		ĺ		MRSELAKFEDIRGVAFDLDGTL
		1		1		VDSAPGLAAAVDMALYALELP
		1				VAGEERVITWIGNGADVLMER
		l				ALTWARQERATORKTMGKPPV
		l				DDDIPAEEQVRILRKLFDRYYG
1		l				EVAEEGTFLFPHVADTLGALQA
1						KGLPLGLVTNKPTPFVAPLLEA
						LDIAKYFSVVIGGDDVQNKKPH
	ŀ	l				PDPLLLVAERMGIAPQQMLFVG
		1				DSRNDIOAAKAAGLDAOARRK
		1				NAGYYYYLRLAKRQMRAGVA
		1				EALRODLLAAKLNVKLVNAGD
						YKFKQIASEKLLIVVTSTQGEGE
						PPEEAVALHKFLFSKKAPKLEN
						TAFAVFSLGDSSYEFFCOSGKD
						FDSKLAELGGERLLDRVDADV
						EYQAAASEWRARVVDALKSRA
1	ŀ					PVAAPSQSVATGAVNEIHTSPY
						SKDAPLVAWRLPNVEGRARAG
		1				GASSFLADRVEEEGEVRVFIEH
				1		NDNFRLPANPETPVIMIGPGTGI
						APFRAFMQQRAADEAPGKNWL
		l				FFGNPHFTEDFLYQVEWQRYV
24386	54754	A	24524	950	2124	MSLISSVIAATWLHVKSYRLSC
24300	34734	1	24324	550	2127	GFYACQSLVLVS/IFPTLSCLFA
		l		i		AEQLLIWSASAFITKVLLVPLIM
		Į				TYAARNIPQNIPEKALFGPAMM
i						ALLAALIVLLCAFVVQPVKLPM
1			1			ATGLKPALAVALGHFLLGLLCI
1		1				VSQRNILRQIFGYCLMENGSHL
1						VLALLAWRAPELVEIGIATDAIF
		i				AVIVMVLLARKIWRTHGTLDL
						TPLLFSLLCFACRKRRLSATRTV
	l	1			1	TVLHSLGITLLLILALWVVQTA
		1			1	ADAGEIFAAGLWLHIDGLGGLF
	İ	1	1		1	LAILGVIGFLTGIYSIGYMRHEV
	1	1			1	AHGELSPVTLCDYYGFFHLFLF
	1	1		1	1	TMLLVVTSNNLIVMWAAIEAT
	1	1	1		1	TLSSAFLVGIYGORSSLEAAWK
						YIIICTVRVAFGLFGTVLVYANA
		1	1			ASVMPQAEMAIFWSE
	<u> </u>				J	MOVIME VACINIAIT WOE

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequen <del>ce</del>	deletion, \≔possible nucleotide insertion)
24387	54755	Α	24525	1	757	RWRLPGVPPFNGFFSKFPLFAP
						GFALSVEYWILLPAMILLMIESV
						ASFAWFIRWFGRVVPGKPSEAV
						ADAAPLPGSMRLVLIVLIVMSLI
						SSVIAATWLHVKSYRLSCGFYA
					1	COSLVLVSIFATLSCLFAAEOLL
		1				IWSASAFITKVLLVPLIMTYAAR
		l				NIPQN\IPEKALFGPAMMALLA
		ı				ALIVLLCAFVVQPVKLPMATGL
					1	KPAL/AGLATDAIFAVIVMVLL
	,	ı				ARH\IWRTHGTLDVNNLHALK
		l				GIMSCSYSS
24388	54756	Α	24526	249	3943	
24389	54757	A	24527	1	1278	
24390	54758	В	24528	48	I44	
24391	54759	В	24529	27	239	
24392	54760	Α	24530	1	551	
24393	54761	Α	24531	3	258	MLFDEPTSALYPEMIN*VLDVM
						VELANEGMTMMVVTHEMGFA
					i	RKEANRVIFMDEGKIVEYSPKD
0.1001			0.500		0.00	AFFDDPKSDRAKDFLAKILH
24394	54762	Α	24532	2	979	
24395 24396	54763 54764	A	24533	1	4116	MLIAEGLMTAKNITVITNSLPA
24390	34704	^	24334	'	1193	AFALSENKDITLVVCGGTVRHK
		١.				TRSMHGSIAERSLODINADLMF
		1	Ì			VGADGIDAVNGITTFNEGYSIS
		l				GAMVTAANKVIAVLDSSKFNR
		l				RGFNOVLPIEKIDIIITDDAVSEV
						DKLALQKTRDTSLVYVLSLADF
		l				FRTASTIGERDGTOVEMILFAGE
						VYLVISLSASLLVLTDCSTEVK
						KGEVVVVCGPSGSGKSTLIKTV
						NGLEPVQQGEITVDGIVVNDKK
		1				TDLAKLRSRVGMVFQHFELFPH
		1				LSIIENLTLAQVKVLKRDKAPA
		l				REKALKLLER/VSGFFAHANKF
		l				PAQLSGGQQQRVAIARALCMD
		1				PIAMLFDEPTSALDPEMINEVLD
		l				VMVELANEGMTMMVVTHEM
		l				GFARKVANRKHTTRSTIALQSL
		l				ATCT
24397	54765	Α	24535	383	962	TDNDRPGLGTDMKKASAIVTN
		l				VAVVLSGRYRRERASGEWAAL
		l	ĺ	1		SFLRPHTLTVFVIPHIQWDDPQ
		1	1	1		MSRP\SGRHLFRCRTVQRKHAL
	l	1	1	1		QVVEKFRPFFPSDRYRYGAPGD
	l		1	I		YAQSFGGQIDDVVKAIQTPSKF
		ĺ				RRFLRVSFD*IRLPILHSGKSPLR
	l	1	ı	1	1	
						RMPNLMRRHSHTARHYPALGT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \=possible nucleotide insertion)
24398	54766	Α	24536	1477	2387	
24399	54767	Α	24537	I	1560	
24400	54768	Α	24538	305	718	
24401	54769	A	24539	1	2161	MNDVDRVGGKNASLGEMITINL SGMGVSVPNGFATTADAFNOF LDQSGWNQRIYELLDKTDIDDV TQLAKAGAQIRQWIIDTFOPEL ENAIREAY AQLSADDENASFAAV RSSATAEDMPDASFAACTLVFG QNLSTSNKDVIELPDNQVDLDK WNVIYPYTFGLIDQLRAKPIMS QANPELEQQIANYEYRIGIGDV LMTVVWDHPDLTTFAGQYRSA SDTGNWVNADGAIFYPYIGILK VAGKTLTQVRNEITARLDSVIE SQVDVSVAAFRISQKAYVITA EVSKSGQQPITNIPLTIMDAINA AGGLTADADWRNVLTQNGV KTKVNLYALMQRGDLRQNKLL HPGDILEIPRNDDLKVFVMGEV KGSTLKMDRSGMTLAEALGN AEGMNQDVADATGIFVIRATQ MKQNGKLANIYQLNAKDASAM ILGTEPQLEPYDIVYTTAPLAR WNENSILKKLERFYEDVKEMPL GEALEQHTGVSFENSKEMPL GEALEGHTGVSFENSKEMPL GEALEGHTGVSFENSKEMPL GEALEGHTGVSFENSKEMPL GEALEGHTGVSFENS
						GSPLSKAADILFPVISDSIRQQA
		L				LPAYSQHISVESTQFSNQGTMA
24402	54770	С	24540	I	1389	
24403	54771	Α	24541	405	576	CEMGAPPWPGCWLS\LFDVDG RFVLAWMLFQGSYLAGALGDQ PVHIHLGQLLALWRPL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24404	54772	A	24542	I	1127	MARTLAHELIDFGEIKRGLLGIK GTEMSADIAKAFNLDVQRGAF VSEVLPGSGSAKAGVKAGDIIT SLNGKPLNSFAELRSRIATTEPG TKVKLGLLRNGKPLEVEVTLDI STSSSASAEMITPALEGATLSDG QLKDGGKGIKIDEVVKGSPAAQ AGLQKDDVIIGVYMDRVNSIAE MRKVLAAKPAIIALQIFPVDHR GAIGGQTTDIPPDPSGINIDRC MKGLSFAGEMTYTNTAHRRA FTNQPRDFMSVAWLGKGIALP SVOWCYKSPVAGRIASLFATLHAI GUESDDGVEULHVGIDTVKLDG QIFSAHVNGEAGLKRVVKKEH ADVQRHAESVYRGQAGQTT DGIRRQDNRVQPGCERCTHQLLR
24405	54773	А	24543	95	599	QIMMLNGSSTPNTSLPTPQAF DRSLPINATSARPSSTSTRHSAE SSISRDCVSNVS/HSGG*TGGIK G*CHAHPADLTACHEADSRGSP SHAYAAVALHVID/GWYLGDQ TAAYATGSAVSCPSSGNQFSRP SAAGRFSAVSAPQGAQRHPGR ARREPRERTAGGCER
24406	54774	A	24544	640	890	LANAQLGSNEISIFQCFAAIQRFI KLHRDASVINHPLAE/TRK*YPV SVYHLRHPPAIIR*PAIRDNA**I LPVTPGYSCSRHQ
24407	54775	А	24545	3	257	ETDIFICDSREFFHHFAKASVHI* PVVSIANFAIQSGKILALGAENI ATTSDRRAQRQKAVCKAHDAS SVSFRTLSVRPTNDCP
24408	54776	A	24546	332	1047	RTRPYCPGMGNEPMTGWMAA AVVTLMIRMCFSVYTMLSESC QRMVIVGYGRCFADRQNLMVC LRSMPNVFTGSCARMRCCLSEN LLYRHRNGHIQAEWP*KKAISD KQYRTSCWERWNAASATIFR/ LSPVEWLTDNGSCYRANETRQ FARMLGLEPKNTAVRSPESNGI AESFVKTIKRDVISIMPKPDGLT AAKNLAEAFEHYNEWHPHSAL GYRSPREYLRQRACNGLSDNR CLEI
24409	54777	Ā	24547	1200	1544	LTGVADGNVDKPCYKFARLW YWIYPAFPPIRRVKVAYREYRP PTTPRVRQVRDLTFCHRPAIPDR MHRY*SSHRALPGCWSKVPAQ KSSRTVSGTD*VDLLSPRAQRA PSPLQG

SEQ ID NO:	nf peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24410	54778	A	24548	263	625	
24411	54779	Α	24549	742	851	VSAQQRVILNKIAVLNRK*RYG CWRLWICPYISRHLLSLNPLQA RCRRYSRGER
24412	54780	A	24550	I	532	MKTLLIIDANLGTIRLYGEDPAG AAARKAKLEIIDNPNDAEMAIV LGDSIPNDSALNGKNVWLGDIS RAVAHPELFI.SERKGHAKPYTA CPTGVAHTFMALKPLKPKKN VAGIPDTGQVCETVQTSCW/AA VETPLRQRSSVVSSGVADG*WF MLPG
24413	54781	В	24551	1	888	
24414	54782	Α	24552	1	2283	
24415	54783	A	24553	992	1662	KKAISDGALTGSSSA VITERDC VSRSRWTAVIRHCTGRSLPAA STVKQYRTSCWERWNAASATI FRLSPVEWLTDNGSCYRANET ROPARMLGLEPKNTAVRSPESN GIAESFVKTIKRDYISIMPKPDG LTAAKNLAEAFEHYNEWHPHS ALGYRSPECYLRIGILVSFISSLQ KFNDFAFEDGLFSGYDAEKRQ YEKSSWNYQLDENGYAKRDET LTHPRC
24416	54784	Α	24554	1	2262	
24417	54785	Α	24555	158	410	
24418	54786	В	24556	1	2607	
24419	54787	A	24557	715	1016	KSAAVPRSGASSQSDSAHFAQP EYAAFSRFPRGSFRHPRAA*YF GRLNHHVRSERGINP*YHRFNL HWRYVADAISADGAFRTVSGG GGRRSGAGRVDWQ
24420	54788	A	24558	652	1214	RTDHCVSRSRWTAVIVRHCTG RSLPAASTVKQYRTSCWDRWN AASATIFR/LSPVEWLTDNGSCY RANETRQFARMLGLEPKNTAV RSPESNGJAESFVKTIKRDYSW PKPDGLTAAKNLAEAFEHYNE WHPPSALGYRSPREVLGQRACT GHAIGSGPSAFFGAPVRRVSRT SVTAKACPRGLL
24421	54789	Α	24559	647	1976	
24422	54790	A	24560	289	574	ETPASSPTLRTSIPSISWMTRGSL RRVMLLCPFPSESGTVSEKAWP EWSSFSSSPPSCRTSASSPPSHLR /NIDVSPKHVVFATIPRNYTMSF LPR
24423	54791	A	24561	816	1375	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nuclcotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ŀ		sequence		
24424	54792	A	24562	li	3993	I IMSOKLKVVTIGGGSSYTPELLE
24424	34/92	A	24302	l'	3993	GFIKRYHELPVSELWLVDVEGG
		1			1	
		l	l			KPKLDIIFDLCQRMIDNAGVPM
		1				KLYKTLDRREALKDADFVTTQ
		1				LRVGQLPARELDERIPLSHGYL
		1				GQETNGAGGLFKGLRTIPVIFDI
		l				VKDVEELCPNAWVINFTNPAG
1						MVTEAVYRHTGFKRFIGVCNIS
1		1	ĺ		\	IGMKILKTRTLDTTGSTSTGDTI
İ						LTQSLLMELSALCRVEVEEGLA
		1		1		LVALIGNDLSKACGVGKEVFG
		l				VLEPFNIRMICYGA
24425	54793	Α	24563	1136	3608	KKAISDGALTGSSSAVITERDC
				1		VSRSRWTAVIVRHCTGRSLPAA
		1			İ	STVKQYRTSCWERWNAASATI
						FR/LSPVEWLTDNGSCYRANET
						RQFARMLGLEPKNTAVRSPESN
					i .	GIAESFVKTIKRDYISIMPKPDG
						LTAAKNLAEAFEHYNEWHPHS
1		1				ALGYRSPREYLRQRACNGLSD
		1				NRCLEIYGQIQLGTGELNITTGV
	ľ					VVTARDTHIALNFOKRRTTGLA
						TOSRORCONRRRVMGKIVIYRE
I					1	AIRFTTQLQTATSINKRTQRVSG
		i				ICRRYTNMASRCNRHOAIVHIM
				1	ł	FAYORPFHFAHFFAIQONFPFGS
1		1			1	IGGQLFRLPVTLLTHQLLLAPA
1		1			l	AHRHGLFQVDVIFRPDNTALPR
1		1			1	NDTHOMVELFLDRFOVVKDIG
	1	1			1	VIELKVVEDORTRAVMNKFRA
		1			i	FVEKGAVILIRFDKKKDIFLQYR
1		1				FHTRISTGHGVANNHQIRLRLE
i					1	LAGIVPLNOLNPLPLNSVPMINI
		1			1	MNGGENADNNVDIQEFMIQPV
		1				GAKTVKEAIRMGSEVFHHLAK
		1			i	VLKAKGMNTAVGDEGGYAPN
		1				LGSNAEALAVIAEAVKAAGYE
1	i					LGKDITLAMDCAASEFYKDGK
		1				YVLAGEGNKAFTSEEFTHFLEE
1	1	1	1			LTKOYPIVSIEDGLDESDWDGF
1		1				,
			l		İ	AYQTKVLGDKIQLVGDDLFVT
	1		1		1	NTKILKEGIEKGIANSILIKFKAQ
			-			HAGLLHMYTCAMLGVGGQCW RYCMSQAFKEDRFAQDQRVAV
24426	54794	В	24564	83	2851	K I CHISQAF KEDIG AQDQK VA V
24426	54795	В	24565	1 I	1578	
24428	54796	A	24566	574	753	
24429	54797	A	24567	1	2112	
24429	54798	В	24568	27	969	
24431	54799	A	24569	1	1773	-
24471	24177	1~	2 1202	1.	r	L

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24432	54800	A	24570	822	984	PSALCGCHSL*CSNASARFFAA VNPSGAFGCRADHPDGELFLTG YVADGGYHS
24433	54801	Α	24571	1	4287	
24434	54802	Α	24572	1	2384	
24435	54803	A	24573	3	3682	ACSARPSEAELELLOTLANADN TIDGILVQI-PAGIDNYKVLER HPDKDVDGFHPYNVGRLCQR APRLRPCTPRGIVTLLERYNIDT FGLNAVVIGASNIVGRPMSMEL LLAGCTTTVTHRFTKNLRHHWE ADALLINAVGRPGFHEDBWIKE GAIVIDVGINRLENGKVVGDVV FEDAAKRASYITPVPGGVGPMT VATLIENTLQAMSNMATFSLGK HPHVELCDLLKLEGWSESGAQ AKIALBGGOVKVD
24436	54804	A	24574		1382	MLRCVNVLADAVKYTLEPKG RNVVLDKSFGAPTITKDGVSVA REIELEDKFENMGAQMVKEVA SKANDAAGDGTTTATVLAQAII TEGLKAVAAGNNPMDLKRAIS RSRKGYGPARIROELNOKGISR EATEKAMRECDIDWCA**RF*S DKHHHRT*RQ/PTRHRYPDESV PGSHNMRSAHALSSAQGYILLP RAVLSGWREENVHPTPGYDSA SGSVEVPPPMTSRRLPAPDNFA APHRYRVLKTGLLSPGCHVRN ADTL
24437	54805	Α	24575	466	1215	
24438	54806	A	24576	1104	1923	LAIPMKDETTEAWLRANHTLISE GAEHIHADSRITPSARELLESR HLRIKFIDEOGRLFYDDEOQOP QPVHGLTSSDEHPQACCELCRQ QPVHGLTSSDEHPQACCELCRQ PVAKKUPDIT.THLSAEKMYAKS DPRLOFRAVLDSTIALAVWLQI ELAFPWQPWLADIRSRLGNIMR ADALGEPLGCQAIVGLSDEDLH RLSHQPPLRYLDHDHLYPEASHG RDAALLNLLRTKVRETETVAA QVFITRSSPEVLRPDILQALNRL SSTYVYMMILSYTKQPLTVKQI QQRLGETQ
24439	54807	A	24577	243	644	HAGSVCPSLYRCQRTAKRSGFY LLLKRLWSFSGNI*IHSGQQNPG RNQQLGLPRSRKGESSRAHPSC QRRPQVWPHPLAGSRPPRPWA ACARLGPIAPARQIPERWPGQV PVLADAHVLPCLAGSLRPALYD F

NO: of peptide hod in USSN location of first codon for last amino a	
sequence 09/540,217 codon for peptide of peptide sequence	deletion, \=possible nucleotide insertion)
24440 54808 A 24578 237 454	RSRLPEGVPVCWCNHSVR*RDC
	RFPVGKLHRLWL/LALPGVRLIF
	RASTKRITGYINARSASIASASA
	RNRLV
24441 54809 A 24579 377 1155	KAYSFATMKSLAISCRWQNA*S
	HVSTFVMVRW/LKGVQFRNHEI
	IGDIVPLAKRYAEEGADELVFY
	DITASSDGRVYPTLITRLADRFG
	VQCIVVGIDTWYDARNLKDISV
	VGLLGAQVPSVRAGMPQSRRQ
	TEGAQGMVRNEEGGSLRLSHH
	QACKATHTQQWTLEVTAQSSY
	NLFPEQWVKLFWGIRQGDLPL
	AFSQKLSRQTFQQQQTQTPRPI
	VGQAPKDLQTGTGTPQTHAVH
	SGDHAAGEQSCKSCKAPIYPLT
	AQFSLETRSFWGSEWGYSTHG
	ELRRRDKGRRRFY
24442 54810 A 24580 1107 1458	RTFWWTTCNSLAPCAGGYPST
	GAALPVSRKLPGRGLLLSGWG
	EHNARYRAGTFLWALFCAAAV
	AYWRRC\WGQYGMWFLRIRRG
	TLGDGTIRRVFVGSLWAAVVIV
24443 54811 A 24581 171 352	EIGVEHIGVFKA PPAPPAESDGLPAHTTRARTTA
24443 54811 A 24581 171 352	DDLRNKRROHVGC*YRTRGSP
	RPNRTWWRVCRYRSPHR
24444 54812 A 24582 5 226	RFINKIWWKVCKIKSFHK
24445 54813 A 24583 476 1939	RPESOHNTAKPAMKVLEOARR
	FLGIRWLSAPASLAN/PVILLVV
	GNILNIVLDVWLVMGLHMNVQ
	GAALATVIAEYATLLIGLLMVR
	KILKLRGISGEMLKTAWRGNFR
	RLLALNRDIMLRSLLLQLCFGAI
	TVLGARLGSDIIAVNAVLMTLL
	TFTAYALDGFAYAVEAHSGQA
	YGARDGSQLLDVWRAACRQSG
	IVALLFSVVYLLAGEHIIALLTS
	LTQIQQLADRYLIWQVILPVVG
	VWCYLLDGMFIGATRATEMRN
	SMAVAAAGFALTLLTLPWLGN
	HALWLALTVFLALRGLSLAAI
	WRRHWRNGRPGTGCLNTKGS
	QQQHRHDSTTQHIHLQLLVIRV
	KTTTTNNSQQTPQYPCQQYHT
	DSRQHRQQNRMVCQRIPHALPS
	PRFRQISRGSTAKVPCAKARAF
	LRVKSDPTLITRLADRFGVQCIV
	VGIDTWYDAETGKYHVNQYTG
	DESRTRVTQWETLDWVQEVQK
	RGAGEIVLAYVSE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24446	54814	Α	24584	1	50	
24447	54815	Α	24585	354	456	HLFSEACCSPGVLRELCRPF*SP
						GTDRNLVMHGH
24448	54816	A	24586	494	835	ITWLPGFRPPFQGSEQFCLAGVP
		1				GATGGSESVLHTRNSYKMLEV
		1				RREVAALISALKVSAKCRNCGS
						SHALTEP/ASSPCWRETSTRPTG APAPTIPPPAASPSPGPMCVOPH
						GED
24449	54817	A	24587	3	726	GED
24450	54818	A	24588	i	1635	
24451	54819	A	24589	1614	2454	RTTHAPASTPNTGTMLCRPYPC
21131	3.017	ľ.	2.507	1.011		CWRRKTP\IIFRATPQWFVSMD
		l				OKGLRAQSLKEIKGVQWIPDW
ŀ		l		1		GQARIESMVANRPDWCISRQRT
						WGVPMSLFVHKDTEELHPRTL
		1		ļ		ELMEEVAKRVEVDGIQAWWD
						LDAKEILGDEADQYVKVPDTL
						DVWFDSGSTHSSVVDVRPEFA
		l				GHAADMYLEGSDQHRGWFMS
		1			l	SLMISTAMKGKAPYRQVLTHG
		l			1	FTVDGQGRKMSKSIGNTVSPQD
					i	VMNKLGADILRLWVASNDYTG
						EIAGSDENLKRACHYNACSISL ARRRH
24452	54820	A	24590	362	517	TKSLISSMSSISSPPVVLICRNSS
24432	34620	^	24390	302	F"'	PEV**RKI*LGTELEFVIMVIAVS
		l				AWP
24453	54821	A	24591	250	930	
24454	54822	A	24592	1388	1912	VCPAAMTGFAGSPFCTAVVTFP
		1			1	DITPLLWILMC*LAILMAESEPK
İ		1				AISPPTISVVP*YAAESLLRVRIP
	ľ	1				VPSFVTSPLPLILCVIAQLRSVE
		1	ì			MVSRPLLITEASPRFFCSLTLSA
		ı	ĺ			LPVSMLNCPTKRLLSPSNSTSVL
		ŀ				AVALTIPFPLMLLRSSPVTFFSV
		_				NRPSWAIFPLPNP
24455	54823	В	24593	1	6018	WALOA BOWLINIERDI DESIRENT
24456	54824	A	24594	510	1003	VALQADQKHKHERPLPPPIPTM FOLAQPGVQLHAKPVYQVP/V
	1				1	MMOATEPTGI*VRIPVLSISTNA
						PSITSIWPLFVRKISVLTLASVV
					1	AASHAASSPRQCAIRCKIWGRW
					1	AGSLRRDDGFGVRSRGSRYGA
					1	SVSISNFSAGINGKSSRSSSPRRS
		1			1	SOTOPVMPI

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24457	54825	A	24595	197	4314	RASTELSLRCCAPSLQ'ALLHFL YLACETWMLLEGI.HLELTVRN LKYANYTSTERFEKREMYPVG YGIPAVIIAVSAIVGPONYGTET HCWLKLDKGFIWSFMGPVAVII LINLYFYFQVLWILRSKLSSLNK EVSTIQOTRYMTEKAISQLEILG CSWGLGFFMVEEVGKTIGSIIA YSFTIINTLQGVLLFVVHCLLNR QINSLLTWTLWTLWTLRQRLSSVN AEVSTLKDTRLLTFKAFAQLFIL GCSWVLGIF
24458	54826	Α	24596	1409	1573	
24459	54827	Α	24597	1283	1563	
24460	54828	A	24598	281	839	CVPTTDL VVQSQI PR WRNWPR CSCSHVSMKAASSPERTFTYCS GIYWLISGSGIGVSVPLLISIGES VINLAVRPVIAHAIRVAITTRQT TRYAVTVVVALPVFRA VEHV FQHLTQTRTGFRWHQFFFNRRQ REOSTTQGEOGRWANFONGKV VNHOLAQRLRRLIGRVRFENQ RFVVFDHLA
2446I	54829	В	24599	I	5652	
24462	54830	A	24600	559	1665	RVERKATIEMKSQRTTSDSWLII LAKQTGIIRAAKKGKTFILHD PROFPYANGSIHIGHSVNKILKDII VKSKGLSGYDSPYVPGWDCHG LPIELK VEQEYGKPGEKFTAAE FRAKCREVAATQVDGQRKDFI RLGVLGDWSHPYLTMDFKTA RIKALGKILONGHLHKGAKEP HWCVDCRSALAEAEVEYYJDK TSPSIDVJAHPFMGFDVPAILGD YVIGQKYGLETANPVGPDGTY LHGTYPTLDGVNVFKANDIVVA LLQEKGALLHVEKMJHSYPCC WRIKKTPIIFRATPGWFVSMDQK GCDIGFHASSNGKRADDSTDD HVRKILCARQRGQPHPHQFQRT

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
24463	54831	A	24601	1	1190	MITDFRESDTMSLFDKKHLVSP ADALPGRNTPMPVATLHAVNG HSMTNVPDGMEIAIFAMGCPW GVERLFWQLPOVYSTAAGVTG GYTPNPTYREVCSGDTGHAEA VRIVYDPSVISYEQLLRKYRCV AVALLGYFKCFPILLNESYKSM QDDLAFIARFIFKDLKFRRFSL KADQKSRIYERVLSIAGVNSWK DKQHQPRLVEHLLVCAESWVA PRALFDAAIEVLAHGKIAIPAYL VAFLVLGPLVLSLIPSQMLPSTL WSSITYSIVVIVEAGVQYHSRLR RFGRVEVNEQIEYLINPIRITRA RAVNFVDNNNRLQAPRILRQDS WWITRPLFGLTWAIIGSPGILPQ VYKSARVMKQAVAVLEPFEA SKEQGKTNGKMVIATVKGDVH DIGKNIYGVVLQCNNYEIVDLG WYMPAEKILR*RLGGIPFTL*R SNRVPGPPEDRISCLFGCVYW SNRVPGPEDRISCLFGCVYW SNRVPGPEDRISCLFGCVYW RCLSSLRRCCRR*RGQAS RQ*LLSSKLAISTILLGLRFGR
				:		VEVNEQIEYLIHNPIR/IARQGG QLC**QQSASGPTDITAGFMVD HQTFTGADVGDNRVARNPATG GQIGARHETGGGLPRTVY*SQQ RAGQNQRQDGDRHREGRRPRH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24464	54832	A	24602	1	1591	MGILVSPLLAPLIGGLLDTMWN
		l				WRACYLFLLVLCAGVTFSMAR
						WMPETRPVDAPRTRLLTSYKTL
		1		·		FGNSGFNCYLLMLIGGLAGIAA
		1				FEACSGVLMGAVLGLSSMTVSI
		l				LFILPIPAAFFGAWFAGRPNKRF
		l				STLMWQSVICCLLAGLLMWIP
						DWFGVMNVWTLLVPAALFFFG
						AGMLFPLATSGAMEPFPFLAGT
						AGALVGGLQNIGSGVLASLSA
						MLPQTGQGSLGLLMTLMGLLI
						VLCWLPLARGYRLSSWTSTTAS
						EDIDCEEGDGEGNETHGLQLAV
				1		EPEGLQPNYMGRGKIRFRDTCL
		İ				LFCTQSRGSQWGFARTELVALL
						PPFLSGAADYRAGLALWLARG
						ADCDVDERHRADRQSNLARSS
						GGFIALAAGAKSDRRLRELNQS
						LQKELARNQHLAERLLETEESV
						RRDVARELHDDIGQTITAIRTQ
						AGIVQRLAADNASVKQSGQLIE
						QLSLGVYDAVRRLLGRLRPRQ
						LDDLTLEQAIRSLMREMÉLEGR
						GIVSHLEW\QSMNQR*AKTSA*
						RCFVSARKG
24465	54833	A	24603	1	1752	
24466	54834	A	24604	1	699	
24467	54835	Α	24605	965	5516	
24468	54836		24606	84	3760	
24469	54837	A	24607	27	278	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
		<u> </u>				
24470	54838	A	24608	311	2529	REWGSTRRCERRISLSRNIARW
						RTIWRITRPCRRCVFRFIASTLR
					. 8	LHLHLIMNDLPLEEYPCVRDLV
		1				PGRFLKFL/HRDPKGTTLVLASI
						LPGRIPVDPRSIIVMLFLAVRWL
						QRQLAGQELLETRATRILNGER
		1	1			GSNVLGTIYEWPPRTSSALDTL
	1	1			\	LREIQNAREQHSRLDTLIRSYA
		1				AQDVKTGLNNRLFFDNQLATL
	ł	I	1			LEDQEKVGTHGIVMMIRLPDFN
						MLSDTWGHSQVEEQFFTLTNL
			1			LSTFMMRYPGALLARYHRSDF
		ŀ				AALLPHRTLKEAESIAVFRALS
						RLPDLWHRSIGTLTEALPOTDK
				l		RQEQIAVALTSYSGLYGLGYGS
						KEIOLTYSGTAGDLISPGLSAIH
						GRCILNTLVFIVKLRWMENTLH
		l				RFGTHHGCSAVRSSHLDSGSPR
		İ				RSSWHMARARNHIHVLEENMA
		1				LLMEQAIANEGLFYRLLYLORS
		l				LTAASSLDDMLMRFHRWARDL
		1				GLAELLVVLPEAKAVGSVAMS
						MLGSDADWGVVLFTSRDASHY
						OOGOGTOLLHEIALMLPELLER
		1				WIERGWEOOVWRYGFLRYVAF
		1				LTGWSAQNELKANPAKGVSAP
		1				
		1			i	KAPRHLPKNIDVDDMNRLLDID
		1				INDPLAVRDLMYGAGLRLSELV
		1				GLDIKHLDLESGEVWVMGKGS
		1				KERRLPIGRNAVAWIEHWLDL
		1	2			RDLFGSERERRQHAGTNVLQT
		<u></u>				ADQRASRARQEGERLHRAAGR
24471	54839	A	24609	1	1749	
24472	54840	Α	24610	1	2911	MNIDAKILNKILANRIQQHIKKL
1		1			1	IHHNQVSFIPGMQGWFNIHKSM
		1				KVIQHINRTKDKNYMIISIDAEK
		l				AFDKIQQPFMLKTLKKLVAGM
i		ı			1	MAVAAPDDPPPPSVSSSKPSTV
1		l			l	APPGRISPSGRALCTSLPSALSSS
	1	1			ĺ	SKFSLLRIFQVPAKLGMVAVAA
						AAGVNAMLRKVAVAAASKPH
	1	1		1		VEIRODGDOFYIKTSTTVRTTEI
		l			l	NFKVGEGFEEETVDGRKCRVR
		1	1		i	PQSHYSVPVSPLGAHGPLLLEE
		Į				PVSPFAACGAPS
24473	54841	A	24611	249	609	QIEVCVRIKPKAGTVLAYS*RSR
[		1				QERPPAPGMTSLSLAAVSSDNP
1		1		I		VSGDEHIRLARASGRFGIGRVA
1		1		1		OPKRIORRSSTPRAATDLGNPG
		1	1	1		HRPCTVRKSALLVVSVCRRHSS
		1		I		RSLRFRSC
	L					KOLIG KOC

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \*=possible nucleotide insertion)
24474	54842	Α	24612	942	1121	
24475	54843	Α	24613	1010	1658	EISSSKRSRTVCKRRAPMFSVDS
			İ			FTCHAASAMRLIPESVNSIFTPS
1				ŀ		VAISALYCTVIEAFGSVRIRSKS
		l			l	LVVSACSSTRIG\DDLAVPGSGQ
		1			i	TVSQSGTRQKR*TECGQF*PSR
						VWWKRCNLLPAAAGHAERLH
1						EKHPDRKTRRVRLVWLAGPPT
ļ		1				RLLGTMGAGTLVLGACCLPWP
		1				REGAHNVATAACGGWRARCD
						LRFRLSCKLGLSRQFFGWNA
24476	54844	A	24614	1	1995	
24477	54845	Α	24615	1	2853	
24478	54846	A	24616	1	1755	
24479	54847	A	24617	2942	3454	SVPDWLMCTAFLSLSRINSPRL
		1				LKAELLLTSSIPVLLPSLLLPRFQ
		1	l			VPSLFATAVPVSAAPLIIAPSV*L
		l				LKTPPADIIRPVTLPLLLNTPAFC
			l		i	TILLMVPVLFTNELSAPAFMVIC
		l			ł	AVISPALLRIPPLTSTLFVITPAF
		ı				LMTAPLSSFTKLLMMPPLVFSR
						RPLDTVI
24480	54848	A	24618	1	2959	MKRYRPREILPWKVPPGNYTPK
		1				IFKIKGTGHNPMSAIENFDAHTP
		l				MMQQYLRLKAQHPEILLFYRM
						GDFYELFYDDAKRASQLLDISL
			1			TKRGASAGEPIPMAGIPYHAVE
					1	NYLAKLVNQGESVAICEQIGDP
						ATSKGCCRSGNRNGGRKIVAVS
						LKKVSGAAKPCGHFRSTRYRV
ì						AYSTIINNKAGIWSAFEGVSAT
						RPNPTPRASPAVRTLSRGTPEPR
						KISTDADNLAQGSLKERLPAWL
21101	44040	<del>ا</del> .	24610	443	1405	WGKKENSKWEVLD
24481	54849 54850	A A	24619 24620	1441	1425 2574	
24482	54850	A B	24620	1	1725	
24484	54851	A	24621	160	390	
				892		
24485	54853	A	24623 24624	1	1183 2493	
24480	54855	A	24625	435	829	ASDVLVSEGCR/YRGASALTST
2448/	J4833	l <sup>A</sup>	24023	1433	027	SVICINTDKALRKIIRLPLNGLR
						KPLNAMTPLPGITWPLCITTEKE
1	1	1		I	i	DLSISDRLSTCIVKFSHPEPEMS
1	1	1	l	1	I	
1		1	l	1	1	VRKFVPPAILLCSVCLRFSCPTR
24486	54056	ļ	24626	762	1962	HQKAPAKWILKSCSAWMR
24488	54856	Α	24626	763	1863	
24489	54857	Α	24627	1423	1712	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24490	54858	A	24628	1	2792	MTYLSAODHELIQVVONGHLO
		1				MCGKSETGNEASPVKCIVLOVL
		1				RGLOYLHRNFIIHSQEOTIPDPSI
						SDQQETPLLAGDMAQRGRRGG
		1			·	FYYVTAYIYVDKKAPVHMRTA
	l					AQGLITLCCQGFG\$LLGYRLGG
						VMMEKMFAYQEPVNGLTFNW
						SGMWTFGAVMIAIIAVLFMIFF
1						RESDNEITAIKVMEDSMSSATT
						LTTPQEQVDSLIMQIAEENGLE
		ļ				VLDQLSQLPEGASAVGESSVRS
						QEDQLSRSFICLMAMM
24491	54859	С	24629	213	345	
24492	54860	A	24630	1	474	MHQTTAKTPHTSYPNNYEEHN
						RRANRSKNYDRIRHTTIILKRTN
						PAQTITNLLKSRFNRYNYVQPR
						SCRSIFIHIPGERDLNLCIAPNPP
	i	l		1		AIFPTSIGYLGPDLNNLPTYAPFI
		ł				TTNLTGPAPTTTHPVGNLGTTG
l						NPEPGGCGILIVWCCVNWACL
		l				WDSVVAAPREPYMPAGLLRHR
	İ			ł		RHYRHRREDLIAVPAARQTDE
						NRRPALCHEYAVGHRKIKTNV
	Į.	ı				EHTEPYILPLITHSQPLNRGAFF
						CPGVRFLVVSVLDFEKRL**RR
		ı		i		WRRSPAGI*GSRGAATTLSQRQ
		1		ŀ		AQLTQHQTIRIPHPPGSGLPVVP
						KFPTGCVVVGAGPVRFVVING
24493	54861	Α	24631	I	924	
24494	54862	В	24632	177	468	
24495	54863	В	24633	592	883	

PCT/US01/08631

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
24496	54864	A	24634	1	1926	MATAKANOEIŌEWLGKYGTAR
24450	34004	ľ	24034	ľ	1520	VKLNVDKDFSLKDSSLEMLYPI
	1	1				YDTPTNMLFTQGAIHRTDDRTQ
						SNIGFGWRHFSGNDWMAAVIT
l		1				FIDHDLSRSHTRIGVGAEYWRD
		1				
						YLKLSANGYIRASGWKKSPDIE
		1				DYQERPANGWDIRAEGYLPAW
	1	1				PQLGASLMYEQYYGDEVGLFG
	l	1				KDKRQKDPHAISAEHSPEFRSE
1		1				ALKLVNASVLLPQPVNSACMN
1		1				HNSTTGAVNSKISRRLLNVNW
		1				RCLPRLHVSNASWQNGMKSWL
		1	i			SSKRPRHTSEAPEMKYVFIEKH
						QAEFSIKAMCRVLRVARSGWY
					ł	TWCQRRTRINIAVVRRADGSRT
						SFVYTSYLAKVNEEWKNNVGT
	1	ŀ				GSTVKWPIGLGGKGNDGIAAF
	l					VQRLPGAIGYVEYAYAKQNNL
		1				AYTKLISA*NEVCLY*KTSG*V
		1				QHQSNVPR/RSGWPAAAGIRGV
	1	1				SGGQG*TLL*YAAQMAPELPSS
l		1		1		TPATWRK*TKSGKTTLV/HGST
		1		1		VKWPIGLGGKGNDGIAAFVQR
		1				LPGAIGYVEYAYAKONNLAYT
1		1				KLISADGKPVSPTEENFANAAK
		1				GADWSKTFAQDLTNQKGEDA
		1				WPITSTTFILIHKDQKKPEQGTE
	1	1				VLKFFDWAYKTGAKQANDLD
		1				YASLPDSVVEQVRAAWKTNIK
	1	1				DSSVNFDMANVFFRVTPGNFL
	l	1				VNREASHORRVFAVRAGVDFI
1		1				YRASSDGLRRCRDRRTRFKRIN
0.1407	54066	<del> -</del>	24625	1387	1620	
24497	54865	Α	24635	1387	1639	ESKPLNLGRGSGFLLTRALSAQ
l	1	1				PLWQSPDQHLTPQAGLRSRL*A
1		1				QRQA*QKTASRVV*PKPLWAL
		┞		ļ		VKATSEDISVVL*ESACQC
24498	54866	Α	24636	1570	2516	IEVAMSIVVKNNIHWVGQRDW
	1	1				EVRDFHGTEYKTLRGSSYNSYF
		1				IREEKNVLIDTVDHKFSREFVQ
		1				NLRNEIDLADIDYIVINHAEEDH
l .		1				AGALTELMAQIPDTPIYCTANAI
		1	*	1		DSINGHHHHPEWNFNVVKTGD
l		1				TLDIGNGKQLIFVETPMLHWPD
				1	1	SMMTYLTGDAVLFNYDAFGQH
				1		YCDEHLFNDEVDQTELFEQCQ
	1	1		1	1	RYYANILTPFSRLVTPKITEILGF
		1	1		1	NLPVDMIATSHGVVWRDNPTO
1		1		1		IVELYLKWAADYQEDRITIFYD
	1					TMVNNTPMMADAIAOGIA\ETG
		1	l			PRLAGENFNV\ARNDKHEDPDL
		1				LSFRSERGA
	L	1	L		<u> </u>	EDI KOLKON

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SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24499	54867	Α	24637	24	363	
24500	54868	A	24638	2	366	RATPRHIIVRFTKVEMKEKMLR AAREKGRVTLKGKPIRLTADLS AETLQARREW*QNLSAFACL*R ILFLLHL*SLVWLDMKCWVENS FL*EC*ILAPTLFWLVGFLPRDP LLV*WASL*G*PDLSLWLPLAF FPSFQLW*I*QLCVLELL
24501	54869	Α	24639	232	292	
24502	54870	Α	24640	4	911	
24503	54871	A	24641	2795	3226	SWWISFLMCCWIWFASILLRIFA SMFIKDIGLKFPFLVVSLPGFGI RMMLAS*NELGRIPSFSIGWNSF RRNGTSSSLYLW*NSAVNPSGP GLFLVGKLLVIASISEPVIGLFTD STSSWFSLGRVYVSRNLSISSRF SSLFA
24504	54872	A	24642	1	3285	
24505	54873	Α	24643	1	4368	
24506	54874	A	24644	200	335	YALLNKDSSPWYPFSVPVLASN TRYLWPLSST**GSLSVWLSPD V
24507	54875	Α	24645	174	263	
24508	54876	Α	24646	1244	1354	

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
į.	sequence	1	09/540,217	codon for peptide	of peptide sequence	delction, \=possible nucleotide insertion)
				sequence		
24509	54877	Α	24647	194	2481	GMFRMNLMEPLTQCINAGHEM
						TKAIAIAOFNDDSPEARKITRR
i		İ				WRIGEAADLVGVSSQAIRDAEK
						AGRLPHPDMEIRGRVEORVGY
1		l				TIEOINHMRDVFGTRLRRAEDV
						FPPVIGVAAHKGGVYKTSVSVH
1		l	İ			LAODLALKGLRVLLVEGNDPO
		1				GTASMYHGWVPDLHIHAEDTL
	ŀ	l				LPFYLGEKDDVTYAIKPTCWPG
		1				LDIIPSCLALHRIETELMGKFDE
1						GKLPTDPHLMLRLAIETVAHDY
		l				DVIVIDSAPNLGIGTINVVCAA
	İ	l				DVLIVPTPAELFDYTSALQFFD
		1				MLRDLLKNVDLKGFEPDVRILL
						TKYSNSNGSQSPWMEEQIRDA
					i	WGSMV\IKNVVRETNEVGKAA
						PMVDSLIARVGVMARGNAITLP
					i	VCGRDVKFTLEVLRGDSVEKTS
						RVWSGNERDQELLTEDALDDLI
				}		PSFLLTGOOTPAFGRRVSGVIEI
		Į.				ADGSRRRKAAALTESDYRVLV
					i	GELDDEOMAALSRLGNDYRPT
		1				SAYERGORYASRLQNEFAGNIS
		l				ALADAENISRKIITRCINTAKLP
		1				KSVVALFSHPGELSARSGDALO
		1			1	KAFTDKEELLKQQASNLHEQK
		i i	l			KAGVIFEAEEVITLLTSVLKTSS
		1				ASRTSLSSRHQFAPGATVLYKG
		l				DKMVLNLDSPQSFSPGAGDHA
İ		l		1	1	NSPQSSSPSTGVHINSPQLSSLS
		l		1		AGVHTISPQSSNPSVGVTALSHP
1		l		l		DLVWVFTLTAHSRPVLVWVFT
24510	54878	Ā	24648	1361	1694	
24511	54879	Α	24649	1	3852	MKLMETLNHCINAGHEMTKAV
		l				AIAOFNDDSPEARKITRRWRIG
		l		l		EAADLVGVSSQAIRDAEKAGR
l		ı				LPHPDMEIRGRVEQRVGYTIEQI
i		1		l		NHMRDVFGTRLRRAEDVFPPVI
		l		l		GVAAHKGGVYKTSVSVHRAQ
		1				DLALKGLRVLLVEGNAPQGTA
		I		l		SVYHGWVPDLHIHAEAALLPF
		I		i		YLGEKDDVTYAIKPTCWPGLDI
		1		l		IPSCLALPRIEPELMGK/FDEGRS
		1	i	l		FMPGPP/HHHPLGRSSO*OVSNS
		I	1	l		LDRAAKGN*KPLCHC
			L		L	

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
24512	54880	IA	24650	1	1720	MEKLLLENDSLSKDVSRLETVF
1						VPPEKHEKEIIALKSNIVELKKQ
						LSELKKKCGEDQEKIHALTSEN
İ	ŀ					TNLKKMMSNOYVPVKTHEEV
1						KMTLNDTLAKTNRELLDVKKK
1		l				FEDINOEFVKIKDKNEILKRNLE
		l				NTONOIKAEYISLAEHEAKMSS
1		l				LSQSMRKVQDSNAEILANYRK
		l				GOEEIVTLHAEIKAOKKELDTIQ
		l				ECIKIMKRAPVIPKHTLNTOPVE
		1				DTSLSTPAAPMVDSLIARVGVM
	İ	l				ARGNAITLPVCGRDVKFTLEVL
		l				RGDSVEKTSRVWSGNERDOEL
		l				LTEDALDDLIPSFLLTGQQTPAF
		1				GRRVSGVIEIADGSRRRKAAAL
		l				TESDYRVLVGELDDEOMAALS
						RLGNDYRPTSAYERGORYATR
						PONEFAGNISAMADAENISRKII
		ŀ				TRCINTAKLPKSVVALFSHPGE
					ŀ	LSARSVIDSAPNLGIGTINVVCA
					ľ	ADVLIVPTPAELFDYTSALQFFD
		1				MLRDLLKNVDLKGFEPDVRILL
1		1				TKYTNINASQSP\WMREKFGMA
		1				WESMVLKNVVRETDEVGKGQI
		l			1	RMRADFV\QAIDQRSSTGAWRN
	1					ALSIWEPVCNEIFDRLIKPRSEIT
24513	54881	A	24651	1094	2271	APFHRPHTKQSLQEQTDVVSIN
		1				EFLKQMAF*AQKLKE/WRTSAA
					l .	SDITPAGFCSCRAIKDAEKPGRL
		ł	İ			PHPD\MEFRDRVDQ\RVVSPFGQ
1					1	II\HMRNVFGPRLPRAKDVFPPV
						IGVAAHKGGVYKTSVSVHLAQ
		1				DLALKGLRVLLVEGN\DP\QGT
		l			ŀ	ASMVSREGDPALHI\YQEAPFLP
		1			Į.	FYLGEKDDVTYAIKPTCCPGLN
i						IIPSCLALPRIETELMGKFDEGK
						LPTDPHLMLRLAIETVAHAHDY
						DVIVIDSAPNLGIGTINVVCAAD
1	l			1		V\LIVLTPVE\MFDYTSARQFFD
						MLRDLLKNA\DLKGFEPEVRIL
	1	1				LTKYSNSNGSQFP\WMEEQIRD
l						AWGSMVLKNVVRETG\EVGKG
				1		\QIRMRT\VFEQAIDQRSSTDTSL
						STPAAPMVDSLIARVGVMASR
24514	54882	Α	24652	I	1415	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24515	54883	A	24653	3008	8165	ILIPFFIGLQNRFTQFRLSETKEIT
						NPYAMRLYESLFQSSKTNGSGI
						VSLKIDWIIERYQLPQSYQRMP
				İ		DFRRRFLQVCVNEINSRTPMRL
						SYIEKKKGRQTTHIAIRDAEKA
						GRLPHPDMEIRGRVEQRVGYTI
					i .	EQINHMRDVFGTRLRRAEDVFP
			1		l	PVIGVAAHKGGVYKTSVSVHL
						AQDLALKGLRVLLVEGNDPQG
						TA\SMYHEWVQNLHIHAEDTLL
					1	PFYLGKKDDVTYAIKPTCWPGL
						DIIPSCLALHRI
24516	54884	A	24654	1	848	
24517	54885	A	24655	2354	3640	NVICNQELVYSSRHQFAPGATV
						LYKGDKMVLNLDRSRVPTEYR
						YGVFLNVLLPATKYFQLHDPSL
						RNMELNYSLASEDI.VLKGLRV
						LLVEGNDPQGTASMYHGWVP
						DLHIHAEDTLLPFYLGEKDDVT
						YAIKPTCWPGLDIIPSCLALHRI
						ETHLMGKFDEGKLPTDPHLML
						RLAIETVAHDYDVIVIDSAPNL
						GIGTINVVCAADVLIVPTPVE\L
						FDYASARQFFHMLRPLLKNVD
						LKGFEPDVRILLTKYSNSNGSQ
						SPWMEEQIRDAWGSMVLKNV
		1				KEFAGNISALADAENISRKIITR
						CINTAKLPKSVVVFFLTPVNYL
		1				PGDALQKAFTDKEELLKQQAS
		1		1	l	NLHEQKKAGVIFEAEEVITLLTS
			ļ.			VLKTSSASRTSLSSRHQFAPGA
		1				TVLYKGDKMVLNLDRSRVPTE
		_				CIEKIEAILKELEKPAP
24518	54886	A	24656	667	870	QIVSSEPKYTDFLSKIPFEQRVR
		1				DVQRNGDHTEGAVR*LSIAFFS
		_				ALLKLLAFFTEEVRAAAVVRFV
24519	54887	В	24657	1	1086	
24520	54888	A	24658	2	250	WRTKSLDIPPLIYPNGIVKNILR
	1				[	HLSQLLNVPITRPFSWILRPF*RP
					i	*RKISTSFIRPLFTFLPA**MLIRS
	1	1		l		SVWQ*KTVSW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24521	54889	A	24659	1	1608	MAETAVINHKKRKNSPRIVQSN
	1007	ľ	-1007	ľ		DLTEAAYSLSRDQKRMLYLFV
		1				DOIRKSDGTLLYMTYRKPDGS
		l				GIVSLKIDWIERYQLPQSYQRM
l		ı				PDFRRRFLQCINAGHEMTKAIA
		i .				AOFNDDSPEARKITRRWRIGEA
		1				ADLLGVSSQAIRDAEKAGPLPH
l	1	İ				PDMENRGRVEORGGYTIEPNN
		1				HMRDGVGTRWRRAEDVF\PPLI
ł		1				G\VAGHKGGR\YKTSVSVHLAQ
		1				DLALKGLRVLLVEGNDPQGTA
		1				SMYHGWVPDLHIHAEDTLLPF
i		l			1	YLGEKDDVTYAIKPTCWPGLDI
		ı			1	IPSCLALHRIETELMGKFDEGKL
	i	1				PTDPHLMLRLAIETVAHDYDVI
		1	- 0			VIDSAPNLGIGTINVVCAADVLI
		1		1		VPTPAELFDYTSALQFFDMLPD
		ı				LVKNRDLKGVEPDMRKORHR
						KAKDLLAQDPTAMTTDELLKE
						ATROPICTAYLEKPMSLECECV
						FCLSCTNSLQKEPQGEGVLCPF
		1				CLEASQKNNIRLNRQLGRLVSH
		1				IKELEPKLKKILKMNFKDAEVP
		1			1	RFSSPKPCSSFFSTCYLSKNCSN
					İ	HPCYSPYPGSGWG
24522	54890	A	24660	1	1128	III C 151 11 G5GWG
24523	54891	A	24661	1035	1208	SLIDRIEALGIFRFAAV*GSAAY
1.025		1		1.000		HGV*YERRRTFLCMLFGYFLSR
						RLESLLLPEVQS
24524	54892	A	24662	1	1316	MEIRGRVEORVGYTIEOINHMR
		1		ľ		DVFGTRLRRAEDVFPPVIGVAA
						HKGGVYKTSVSVHLAQDLALK
				i		GLRVLLVEGNDPQGTASMYHG
					İ	WVPDLHIHAEDTLLPFYLGEKD
					İ	DVTYAIKPTCWPGLDIIPSCLAL
				l		HRIETELMGKFDEGKLPTDPHL
						MLRLAIETVAHDYDVIVIDSAP
						NLGIGTINVALQNFQRYTGIQH
-				l		VHRIGMAERMWCDRNRERHT
		1		l		VSSSGGNRLPNPGPDRSDLALK
1				1		GLRVLLVEGNDPQGTASMYHG
		1				WVPDLHIHAEDTLLPFYLGEKD
			1			DVTYAIKPTCWPGLDIIPSCLAL
			1			HRIETELMGKFDEGKLPTDPHL
	1	1		I		MLRLAIETVAHDYDVIVIDSAP
1		1	1	1		NLVFRICFVIWAQERIDIKGFEP
1		1		1		DVRILLTKYSNSNGSQSPWMEE
		l				QIRHAWGSMVLKN/VVRETDE
	1					VGKDTGVRSKSIWCHRNSRWE
	1	1	i .	1	L	A OVDIO A VSVSI M CHK NSK ME

SEQ ID	SEQ ID NO:	Met	SEQ tD NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24525	54893	Α	24663	504	1254	SYVKYFPHQPAQKYFQQIHSAI
		l				GLHNADHVHKHLLGDNRYPIW
		l				IMQPSAHHPARQPEHDIAFGKC
		1				SSSVSVHLAQDLALKGLRVLLV
		1				EGNDPQGTASMYHGWVPDLHI
		1				HAEDTLLPFY\DPHLMLRLAIET
	ļ	l			l	VAHDYDVIVIDSAPNLGIGTINV
	1					VCAADVLIVPTPAELFDYTSAL
		İ				QFFDMLRDLLKNVDLKGFEPD
						VRILVVRVSQRMVVAGCIIAFID
						KLLNVLICFPFSVKNRFVIFHSV
						DFECVIRF
24526	54894	Α	24664	1	3216	MKLMETLNQCINAGHEMTKAI
	1					AIAQFNDDSPEARKITRRWRIG
				I		EAADLVGVSSQAIRDAEKAGR
	1	l				LPHPDMEIRGRVEQRVGYTIEQI
				İ		NHMRDVFGTRLRRAEDVFPPG
		l			1	FGDFADFGTTIKQDFRLLGQTS
						VDRLLQLSQGQAVKGNQLLPV
						SLVKRKTTLAPNTQTASPRALA
						DSLMQLARQVSRL/VKRAPTC
		l				WPGLDIIPSC\LALPRIETELMGK
						FDEGKLPTDPHLMLRLAIETVA
						HDYDVIVIDSAPNLG
24527	54895	A	24665	1	419	MSSSDKDFDFSVANMVNFFAP
						VFPMGKYYTQADKVLIPLAIQE
İ					ŀ	HDGICEIHVAKYAEIFGLTSAEA
	1				I	SKDIRQALKSFAGKEVVFYRPE/
		1			İ	VGCRR*KRL*IFSLVYQTCAQSI
				I	1	QRALQCTYQPISHSLLYRVTEP
						VYAVSA
24528	54896	Α	24666	1	2598	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24529	54897	Α	24667	1	2027	MRGHREKAAACKPRTRASEGT
						TPASTFILDFOPPNCNYTSGDNH
l		1				TLRDPHYVEDKGHKYLVFEAN
		1				TGTENGYQGEESLFNKAYYGG
		1				GTNFFRKESQKLQQSAKKRDA
1						ELANGALGIIELNNDYTLKKVM
						KPLITSNTVTDEIERANVFKMN
1						GKWYLFTDSRGSKMTIDGINSN
	1					DIYMLGYVSNSLTGPYKPLNKT
						GLVLOMGLDPNDVTFTYSHFA
						VPOAKGILCEEDNYTAGDNHM
						MRAPHCEEDRAHKFVVFDANA
1						GTESGHOGESSLFNRACGGGGT
1						FFFSKESQKLQQSAKKRDCINA
i		ŀ				GHEMTKAIAIAQFNDDSPEARK
						ITRRWRIGEAADLVGVSSQAIR
						DAEKAGRLPHPDMEIRGRVEO
ļ.						RVGYTIEQINHMRDVFGTRLKR
						AEDVFPPVIGVAAHKGGVYKT
						SVSVHLAQDLALKGSLLLPKND
						FLFKLGLEGHLPLGKIHSPPTGA
						DVARGSSGLPKSELFLPERNTO
						ELQQDSEEGPLALQVLQSNLM
1						DFADFGTTIKODFRLLGOTSVD
1		l				RLLQLSQGQAVKGNQLLPVSL
1		1				VKRKTTLAPNTQTASPRALADS
		1				LMQLARQVSRLESGQNNDGIC
		1				EIHVAKYVEIFGLTSAEASKDIR
ŀ						QALKSFAGKLVVFYRPE/VGCR
						R*KRL*IFSLVYOTCAOSIORAL
	-	1				QCTYQPISHSLLYRVTEPVYAV
24530	54898	A	24668	1	989	PWISAPVPVDVVEGAMDSVTV
		1				LSFGGLMLYFCAGWPPARRWC
						FPESISCGSMERDQWWGLQVA
						KRAGLAGGQSGRTVLRERVRIE
		1				IASTHIALAARHSDWRCCRNGR
		1				YPARGPAALONFORYTGIOHV
		1				HRIGMAERMWCDRNRERHTVS
		1				SSGGNRLPNPGPDRSCDNLKTC
1		1				HTSHGSVMAETAVINHKKRKN
		1				SPRIVQSNDLTEAAYSLSRDQK
		1	1			RMLYLFVDQIRKSDGTLQEHD
		1				GICEIHVAKYAEIFGLTSAEASK
		ı				DIRQALKSFAGKEVVFYRPE/V
1		1	1			GCRR*KRL*IFSLVYQTCAQSIQ
		1				RALOCTYOPISHSLLYRVTEPV
		ــــــــــــــــــــــــــــــــــــــ	L			

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codun for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
10.	sequence	liou.	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24531	54899	A	24669	350	1203	LLTARLTWVSARLMSDFADFG
						TTIKQDFRLLGQTSVDRLLQLS
						QGQAVKGNQLLPVSLVKRKTT
						LAPNTQTASPRALADSLMQLA
						RQD/LALKGLRVLLVEGNDPQG
						TASMYHGWVPDLHIHAEDTLL
l						PFYLGEKDDVTYAIKPTCWPGL
		İ				DIIPSCLALHRIETELMGKFDEG
						KLPTDPHLMLRLAIETVAHDYD
		l				VIVIDSAPNLGIGTINVVCAADV
		1			ĺ	LIVPTPAELFDYTSALQFFDMLR
					į .	DLLKNVDLKGFEPDVRILLTKY
						SNSNGSQSPWMEEQIRDAEGSR
24532	54900	Α	24670	1	936	
24533	54901	Α	24671	1070	1324	LGTPGGCSRMTCALVPLIPNDD
1	1	ı				TPTRRTRPLLGQGVFSVRS*TVP
	į.	l				ADQSTSGDGDSTCNVRGSCPCR
		_				IAITILITPATPAAACV
24534	54902	Α	24672	1	1212	
24535	54903	A	24673	2373	3381	FKRGQYGQLLRPRFHHGQILYA
ŀ		1	1			RRQGADAAGDSGIW/HGIWTEP
	1	1		l		MTM*KKDER*PSPTQAS*DNW
i		l				KSIRGSCWQALSITRN/CKNSPRI
		i		1		VQSNDLTEAAYSLSRDQKRML
		l			1	YLFVDQIRKSDGTLQEHDGICEI
		l		l		HVAKYAEIFGLTSAEASKDIRQ ALKSFAGKEVVFYRPEEDAGD
		l				EKGYESFPWFIKRAHSPSRGLY
1		l				SVHINPYLIPFFIGLONRFTOFRL
		l				SETKEITNPYAMRLYESLCQYR
		l			i	KPDGSGIVSLKIDWIERYQLPQ
		1				SYORMPDFRRRFLOHIFVLRER
		1		l		PETVLIDLIORTKDAVRELDNL
		1		l		QYRKMKKLLFQEAHNGPAVEA
						OEEEEDD
24536	54904	A	24674	31	811	RRGLLGTRLRLSEDVFPPVLGV
2 1554	1,00	1			l	AAHKGGVDKTSVSVHLAODLA
	1	1		l		LKGLRVLLVEGNDPQGTASMY
		1				HGWVPELHIHAEDTLLPFYLGE
		1			l .	KDDVTYAIKPTCWPGLDIIPSCL
		1				ALHRIETELMGKFDEGKLPTDP
		ı			l .	HLMLRLAIETVAHDYDVIVIDS
1		1				APNLGIGTINVVCAADVLIVPTP
				l		AELFDYTSALQFFDMLRDLLKN
						VDLKGFEPDVRILLTKYSNSNG
				I		SQSPWMEEQIRDAWG\SMVLK
				1	1	NVVAFQNLEFLFILLRIWL
24537	54905	Α	24675	1004	1105	-
24331	154705	111	12-7073	1.00-	1	L

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	1	09/540,217	codon for peptide sequence	or pepude sequence	deletion, (-possible nucleotide insertion)
		1				
24538	54906	Α	24676	1	1139	MEKKITGYTTVDISQWHRKEHF
						EAFQSVAQCTYNQTVQLDIKK
		l			[	KLKTVKKNKHKFYPAFIHILAR
		l			İ	LMNAHPEFRMAMKDVAEYNG
		l				YHVVFALAGSPKDADDTSIYM
						FYQKVGDNSIDSWKNAGRVFK
		l				DSDKFDANDPILKDQTQEWSGS
1						ATFTSDGKIRLFYTDYSGKHYG
			i			KQSLTTAQVNVSKSDDTLKING
1						VEDHKTIFDGDGKTYQNVQQFI
						DEGNYTSGDNHTLRDPHYVED
	1	1				KGHKYLVFEANTGTENGYQGE
		l				ESLFNKAYYGGGTASSIRKSDG
						TLQEHDGICEIHVAKYAEIFGLT
1					ŀ	SAEASKDIRQALKSFAGKEVVF
						YRPE/VGCRR*KRL*IFSLVYQT
ļ			1			CAQSIQRALQCTYQPISHSLLYR
		ļ. —		ļ. —	000	VTEPVYAVSA
24539	54907 54908	A	24677	1	930 2486	MNMNIKKIVKOATVLTFTTALL
24540	34908	l <sup>A</sup>	24078	1	2480	AGGATQAFAKENNQKAYKETY
i			ì			GVSHITRHDMLQIPKQQQNEKY
	i					QVPQFDQSTIKNIESAKGLDVW
ļ	l	I				DKWPLANADGTVAEYNGYNV
l						VDKFDANDPILKDOTQEWSGS
l						ATFTCDGKIRLFYTDYSGKHYG
	l	1				KQSLTTAQVNVLKSDDTLKING
		ĺ				VEDHKAIFDGDGKTYONVOOL
	1					IDEGNYTSGDNHTLRDPHYVED
	l					KGHKYLVFEANTGTENGYQGE
1				1		ESLFNKAYYGGGTNFFRKESQK
						LQQSAKKRDAELANGALGIIEL
l					1	NNDYTLKKVMKPLITSNTVTDE
			ľ			IERANVFKMNGKWYLFTDSRG
						SKMTIDGINSNDIYMLGYVSNS
			l			LTGPYKPLNKTGLVLQMGLDP
						NDVTFTYSHFAVPQAKGNNVV
						ITSYMTNRGFFEDKKATFAPSF
						LMNIKGNKTSVVKNSILEQGQL
	1		l			TWLQVAKRAGLGGGQSGRTVL
						RERLPNYKNFKGTIQELGQNQY
1	İ					AVSGEIFVVDRNTVEITELPVRT
1	1		1			WTQVYKEQVLEPMLNGTDKTP
1						ALISDYKEYHTDTTVKFVVKM
į.			1			TEEKLAQAEAAGLHKVFKLQT
İ			1			TLTCNSMVLFDHMGCLKKYET
1						VQDILKEFFDLRLSYYGLRKEW
i					1	LVGMLGAESTKLNNQARFILEK
1	1	1	1			IQGKITIENRSKKDLIQMLVQRG
1	1		l			YESDPVKAWKEAQEKCDNLKT
1	1	1	l	1	i	CHTSHGSVMAETAVINHKKRK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	į.	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24541	54909	A	24679	5	253	I SAPKVWRGLROOGAGISVLP*V
		1		ľ		CAQP/VQVAKAPELILNFDLPK
	Ì	1				KYFDKL/R/DTHFFSPHTGKFPT
		1				VKILMAPFANSASRGVALC
24542	54910	A	24680	334	842	PDKEGNEIWVDMYTVKPSGWT
	34710	Γ.	2.000	1		VRTFDKPRKRFIAFFIAGILFRAI
		1				KNHFLPRETLQCLPYILTGFRRG
		1				OSEYFSIFSNMDLADTVMFLVG
		1				GHKIFVFHCHTELRMSIHQAGK
		1				NVNKGRIKLVLIFLYGL*/QRP*
	ľ					YPAERSGYRYIEQLTEMPONVL
1						YDAIGIYORWYIO
24543	54911	A	24681	676	1169	DNLKTCHTSHGSVMAETAVIN
24343	34911	Ι^	24001	070	1109	HKKRKNSPRIVQSNDLTEAAYS
						LSRDOKRMLYLFVDOIRKSDGT
		1				LOEHDGICEIHVAKYAEIFGLTS
l		1		ļ		AEASKDIRQALKSFAGKEVVFY
ŀ		1				RPE/VGCRR*KRL*IFSLVYQTC
1		1			ì	AQSIQRALQCTYQPISHSLLYRV
1	ì	1				TEPVYAVSA
24544	54912	A	24682	1	783	TEL VIA VOA
24545	54913	A	24683	i	1884	
24546	54914	A	24684	1	1131	
24547	54915	A	24685	1	2856	
24548	54916	A	24686	528	725	
24549	54917	Α	24687	1	898	
24550	54918	Α	24688	1879	2343	
24551	54919	A	24689	1	1254	
24552	54920	Α	24690	1	1113	
24553	54921	В	24691	1	1156	
24554	54922	В	24692	1	1980	
24555	54923	В	24693	1	1932	
24556	54924	Α	24694	1	663	
24557	54925	A	24695	1123	1764	
24558	54926	Α	24696	2	1772	MELKTKARELREECRSLRSRCN
24559	54927	A	24697	1	1040	QLEERVSAMEDEMNEMKREG
	ľ					KFREKRIKRNEOSLOEIWDYVK
1		1				RPNLRLIGVPESDVENGTKLEN
	i					TLQDIIQENFPNLARQANVQIQE
ł						IQRTPQRYSSGRETPRHIIVRFT
I		1				KVEMKEKMLRAAERKCDNLK
1	1	1				
1	1	1				TCHTSHGSVMAETAVINHKKR
		1		1		KNSPRIVQSNDLTEAAYSLSRD
1		1				QKRMLYLFVDQIRKSDGTLQE
1		1				HDGICEIHVAKYA EIFGLTSAEA
1	1	1				SKDIRQALKSFAGKEVVFYRPE
		1				EDAGDEKSYEQPSKDIRQALKS
				1		FAGKEVVFYRPE/VGCRR*KRL
		1			1	*IFSLVYQTCAQSIQRALQCTYQ
		1	L	L	L	PISHSLLYRVTEPVYAVSA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24560	54928	A	24698	1683	4289	RVFKDSDKFDANDPILKDQTQE
						WSGSATFTSDGKIRLFYTDYSG
		ı				KHYGKQSLTTAQVNVSKSDDT
		1				LKINGVEDHKTIFDGDGKTYQN
	1	1			1	VQQFIDEGNYTSGDNHTLRDPF
		1				YVEDKGHKYLVFEANTGTENG
	į.			1		YQGEESLFNKAYYGGGTNFFR
				ı	l	KESQKLQQSAKKRDAELANGA
				1	i	LGIIELNNDYTLKKVMKPLITSN
						TVTDEIERANVFKMNGKWHSK
				1		ILVSGTHRARTEVEVTDPIHQQ
		1		1		RMGCCFTVGAGEICEKPTGCFF
	1	1		1		TPLPNCNCQWTGSAILDRRSHS
				1		HPHSPASGSWVKVVTGSYQLT
				1		HSPGIGLRDFADFGTPHQTGFSF
		ı				AGANQRGPLVATLSGPGGEGQ
				1		SAVARLTGEKKNHPGAQYANR
- 1		1				LSPRVGRFINAAGTTGFPTGKR
		1				AVSATQLMHFKILVSRAHRART
						ELEVTDPGSSAENGLLFPVVLL
						LRKARKTLMPPHSTCFIKRSAP
	1			1		TQSTAGKTRPRVLKDSDKFDA
		1				NDPILKDQTQEWSGSAPFTSDG KIRLFYTDYSGKHYGKOSLTTA
						QVNVSKSDDTLKINGVEDHKTI
				1		FDGDGKTYONVOOFIDEGNYT
		l		1		SGDNHTLRDPHYVEDKGHKYL
		1	ľ			VFEANTGTENGYQGEESLFNKS
		1	ļ			VLRRRHELLPDFADFGPTIKQD
		ŀ		1		FRLLGQTSVDRLLQLSQGQAV
		i	ŀ	1		KGNOLLPVSLVKRKTTLAPNTO
						TASPRALADSLMQLARQVSRLE
24561	54929	В	24699	1	2034	
24562	54930	Α	24700	1	1038	
24563	54931	Α	24701	1	1149	Was out a complete to the comp
24564	54932	Α	24702	2	823	WPLQNADGTVAEYNGYHVVF
	i					ALAGSPKDADDTSIYMFYQKV
	ĺ	1				GDNSIDSWKNAGRVFKDSDKF
		1				DANDPILKDQTQEWSGSATFTS
		1	ļ			DGKIRLFYTDYSGKHYGKQSLT
		1		i		TAQCDNLKTCHTSHGSVMAET
		1	l			AVINHKKRKNSPRIVQSNDLTE AAYSLSRDQKRMLYLFVDQIR
	1	ı				KSDGTLQEHDGICEIHVAKYAE
		ı				IFGLTSAEASKDIROALKSFAGK
	1	1				EVVFYRPEEDAGDEK/ML*IFSL
		ı	1			VYQTCAQSIQRALQCTYQPISH
		1	1			SLLYRVTEPGYAVSA
24565	54933	В	24703	1	1206	
24566	54934	Α	24704	2	1037	
24567	54935	В	24705	I	1389	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hed	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24568	54936	A	24706	1	602	I MGTGDFICISMTGGAPWGFRLO
24308	34936	ΙΑ.	24700	'	002	GGKEQKQPLQVAKWANLKPGP
İ		1	ŀ			PSHGSVMAETAVINHKKRKNSP
		1			i	RIVOSNOLTEAAYSLSROQKRM
ŀ						LYLFVDQIRKSDGTLQEHDGIC
			İ			EIHVAKYAEIFGLTSAEASKDIR
		1				QALKSFAGKEVVFYRPE/VGCR
						R*KRL*IFSLVYQTCAQSIQRAL
		l				QCTYQPISHSLLYRVTEPVYAV
24569	54937	A	24707	1	774	QCT QTISHSEET RETTER TTAT
24570	54938	A	24708	1479	1947	
24571	54939	A	24709	1	696	
24572	54940	A	24710	1	1773	
24573	54941	A	24711	1	1509	
24574	54942	A	24712	1	891	
24575	54943	A	24713	156	242	VAGRGIGYRLSPNKGL*TWSAL CKPIAE
24576	54944	Α	24714	1441	1866	
24577	54945	Α	24715	1	1098	
24578	54946	Α	24716	Ī	1989	
24579	54947	В	24717	126	848	
24580	54948	В	24718	58	688	
24581	54949	Α	24719	]	542	
24582	54950	Α	24720	1	675	
24583	54951	Α	24721	1	1209	MKMPEKHDLLAAILAAKEQGI
				l	ŀ	GAILAFAMAYLRGRYNGVLEA
}			i			RSLKSRCQQSHALAEGSRGGPP
					İ	VTGRFKSTFTADSWNRYSLPSL
		1				TGGGNILHSRTHRRTVSRLLGR
		ĺ				RWQELLRDVFGTRLRRAEDVF
		l				PPVIGVAAHKGGVYKTSVSVH
		l				LAQDLALKGLRVLLVEGNDPQ
		l				GTASMYHGWVPDLHIHAEDTL
		1				LPFYLGEKDDVTYAIKPTCWPA
		l			İ	LDIIPSCLALHRIETELMGKFDE
		1				GKLPTDPHLMLRLAIETVAHDY
		1				DVIVIDSAPNLGIGTINVVCAAD
		1				VLIVPTPAELFDYTSALQFFDM
						LRDLLKNVDLKGFEPDVRILLT/
					İ	KYSNSNGSQSPWMEEQ\IRDAW
						GSMGLKNV/VRETDEIG\KGQIR
						MRTVFETAMDKRLLTGAWEK
						AFFFLEPGFEEKSNR

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide scquence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24584	54952	A	24722		743	MIGMELPGLEGTELEGCHEREY VYDRWAVPEKRGEARHVEARS PGGGSEKRGWGGARHQRMCR KGADNGGAGRARFARLRQL ASRAIGRRKGAVLYQTPVINHK KRKNSPRIVQSNDLTEAAYSLS ROOKRNLTLYFVDQIRKSDGTL QEHDGICEHVAKYAEIFGLTL WREASKDIRQALKSFAGKEVV TCAQSIQRALQCTYQPISHSLLY RVTEPVYAVG
24585	54953	A	24723	1157	1471	
24586	54954		24724	1	824	MEENERDLEINMINMSKELKVE NIKKHIGMLEKDEIEIVYKSILI EIEIPTQCLRNLNTLHLAASAGN HMSNTACWYGCEDFTSVSTFA HQGVGSWDAYEPAASLLSCD NLKTCHTSHGSVMAETAVINH KKRKNSPRIVQSNDLTEAAYSL SKDQKRMLYLEVDQIRKSDGTL QEHDGICEIHVANYAEIYGLTS ACASKDIRQALKSFAGKEVVFY RPEVYGCRR*KRL*IFSLVYQTC AQSIQRALQCTYQPISHSLLYRV TEPUYAVSA
24587	54955	A	24725	996	1169	SLIDRIEALGIFRFAAV*GSAAY HGV*YERRRTFLCMLFGYFLSR RLESLLLPEVQS

SEO ID	ISEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
i	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
	1	_	24726		1743	DSPE\ARKITERWRIGEAADLV
24588	54956	Α	24726	3	1743	
						GVSSQAIRDAEKAGRLPHPDME
		l			1	IRGRVEQRVGYTIEQINHMRDV
1						FGTRLRRAEDVFPPVIGVAAHK
						GNDPQGTASMYHGWVPDLHIH
						AEDTLLPFYLGEKDDVTYAIKP
						TCWPGLDIIPSCLALHRIETELM
	İ					GKFDEAQPNLGIGTINVVCAAD
1					ł	VLIVPTPAELFDYTSALQFFDM
	l					LRDLLKNVDLKGFEPDVRILLT
ł						KYSNSNGSQSPWMEEQIRDAW
İ					ì	GSMVLKNVVRETDEVGKGQIR
l	1					MRTVFEQAIDQRSSTDTSLSTP
						AAPMVDSLIARVGVMARGNAI
1						TLPVCGRDVKFTLEVLRGDSVE
	1					KTSRVWSGNERDQELLTEDAL
l						DDLIPSFLLTGQQTPAFGRRVSG
	ĺ					VIEIADGSRRRKAAALTESDYR
ŀ						VLVGELDDEOMAALSRLGNDY
l	ļ					RPTSAYERGORYASRLONEFAG
ł						KYFLRWLMRKIFHIITRCINTAK
İ	l					LPKSVVALFSHPGELSARSGDA
	1					LQKAFTDKEELLKQQASNLHE
ł	1					OKKAGVIFEAEEVITLLTSVLKT
						SSASRTSLSSRHOFAPGATVLY
						KGDKMVLNLDRSRVPTECIEKI
						EAILKELEKPAP
24589	54957	A	24727	1	1623	
24590	54958	A	24728	1	1863	
24591	54959	A	24729	i -	2622	
24592	54960	A	24730	ī	1338	
27372	124700	<u>'`</u>	24730	1.	11000	L

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24593	54961	A	24731	1	1823	MNMNIKKIVKOATVLTFTTALL
						AGGATQAFAKENNQKAYKETY
		l		l		GVSHITRHDMLQIPKQQQNEKY
						QVPQFDQSTIKNIESAKGLDVW
						DSWPLQNADGTVAEYNGYHV
		1				VFALAGSPKDADDTSIYMFYQ
						KVGDNSIDSWKNAGRVFKDSD
				1		KFDANDPILKDQTQEWSGSATF
						TSDGKIRLFYTDYSGKHYGKQS
	İ	1				LTTAQVNVSKSDDTLKINGVED
		1				HKTIFDGDGKTYQNVQQFIDEG
		1				NYTSGDNHTLRDPHYVEDKGH
1	ļ	1				KYLVFEANTGTENGYQGEESLF
		1		1		NKAYYGGGTNFFRKESQKLQQ
		1				SAKKRDAELANGALGIIELNND
		l				YTLKKVMKPLITSNTVFIRDKL
		l				MERRNRRTGRTEKARIWEVTD
		ı				RTVRTWIGEAVAAAAAADGVTF
		l	l	ĺ		SVPVTPHTFRHSYAMHMLYAG
		l		1		IPLKVLQSLMGHKSISSTEVYTK
		l				VFALDVAARHRCDNLKTCHTS
		l				HGSVMAETAVINHKKRKNSPRI
		1				VQSNDLTEAAYSLSRDQKRML
		l				YLFVDQIRKSDGTLQEHDGICEI
		l				HVAKYAEIFGLTSAEASKDIRQ
		1	1			ALKSFAGKEVVFYRPE/VGCRR
						*KRL*IFSLVYQTCAQSIQRALQ
21501		ļ				CTYQPISHSLLYRVTEPVYAVS
24594	54962	Α	24732	1476	2221	

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24595	54963	A	24733	1	815	MRVGRKAVGAIWWSSKQELFQ
						QLFLRTELEDIGPGIVKRELGRE
						GSKDTGSSDDWEQLPDWKTKR
						VGNERDQELLIEDALDDLIPSFL
						LTGQQTPAFGRRVSGVIEIADG
			ŀ			SRRRKAAALTESDYRVLVGEL
		l				DDEQMAALSRLGNDYRPTSAY
						ERGQRYASRLQNEFAGNISALA
						DAENISRKIITRCINTAKLPKSV
						VALFSHPGELSARSGDALQKAF
			i			TDKEELLKQQASNLHEQKKAG
						VIFEAEEVITLLTSVLKTVQSKN
						LSHFTWICHAETAVINHKKRKN
			ŀ			SPRIVQSNDLTEAAYSLSRDQK
						RMLYLFVDQIRKSDGTLQEHD
						GICEIHVAKYAEIFGLTSAEASK
						DIRQALKSFAGKEVVFYRPEED
						AGDEKGYESFPWFIKHWVTISP
	1					QTSAYERGQRYASRCKNEFAG
						NISALADAENISRKIITRCINTAK
						LPKSVVALFSHPGELSARSGDA
						LQKAFTDKEELLKQQASNLHE
					i	QKKAGVIFEAEEVITLLTSVLKT
						SSASRTSLSSRHQFAPGATVLY
						KGDKMVLNLDRSRVPTECKRK
						LRPF*SGQRTDTAYVFDPGRDY

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \=possible nucleotide insertion)
24596	54964	Α	24734	2	1249	DFADFGTTIKQDFRLLGQTSVD RLLQLSQQAVKOQLLPYSL VKKKTTLAPNTQTASPRALADS LMQLARQVSRLESGQLAISGRI LLPMLNSINIKLAAPSQCTEPF FRRDPTLHPKSEVRLPPKVTTIT VISNPILITPQEKSVAQCTYNQT VOLDITAFLETVKKNKHKFYPA FIHILARLMNAHPEFRMAMKD GELVIWDSVHPCYTVFHEGTET FSSLWSEYHDDFRQFLHIYSQD VACYGENLAYPPKGTETF FSSLWSEYHDDFRQFLHIYSQD VACYGENLAYPPKGTETF FSSLWSEYHDDFRQFLHIYSQD VACYGENLAYPPKGTETF FSSLWSEYHDDFRQFLHIYSQD VACYGENLAYPPKGTEMFMV AQUENLAYPTMGKYYTQGDKVLMPL AQUENLAYPTMGKYYTQGDKVLMPL AQUENLAYPTMGKYYTQGDKVLMPL AQUENLAYPTMGKYYTQGDKVLMPL SSPFWFIKRAHSPSRGLYSVHIN PYLIPFFIGLONRFTQFRLSETKE ESFFWFIKRAHSPSRGLYSVHIN PYLIPFFIGLONRFTQFRLSETKE LSYIEKKKGRQTTHIIRISGTL QEHDGIGEHVAKYAGFIGLTSA EASKDIRQALKSFAGKEVVFYR EEDAGDERGYESFPWFIKRAH LSYIEKKKGRQTTHIRISGTL QEHDGIGEHVAKYAGFIGLTSA EASKDIRQALKSFAGKEVVFYR PEEDAGDERGYESFPWFIKRAH SPSRGLYSVHINPYLIPFFIGLON RFTQFRLSETKEITNPYAMRLY ESSCLYSVHINPYLIPFFIGLON RFTQFRLSETKEITNPYAMRLY AKYAEIFGLITSAEASKDIRQAL KSFAGKEVYFYRPEVGCR**K A*1*SILYYOTCAOSIORALOCT

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	tocation of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	or peptide sequence	deterion, (-possible nucleonide insertion)
24597	54965	A	24735	1	2286	MLTLILLDLOFMAOLAGYOMS
		l				FNSTKSKQTKNNFAVGYRTGD
						FOLHTNVNDGAEFGGSVYOKV
						CEELDALINLAWTSSTSCTRFG
	İ					LAAKFOLKPIAFISTKVNHWLT
						GVSYTPPLEAGVKLNLSALCDN
1						LKTCHTSHGSVMAETAVINHK
						KRKNSPRIVQSNDLTEAAYSLS
	i				1	RDQKRMLYLFVDQIRKSDGTL
1					i	QEHDGICEIHVAKYAEIFGLTSA
1						EASKDIRQALKSFAGKEVVFYR
						PEEDAGDEKGYESFPWFIKRAH
						SPSRGLYSVHINPYLIPFFIGLQN
ĺ						RFTQFRLSETKEITNPYAMRLY
l						EIPVVSIRKPGMGSGIVSLKIDW
l			ŀ			HERYQLPQSYQRMPDFRRRFLQ
						CINAGHEMTKAIAIAQFNDDSP
						EARKITRRWRIGEAADLLGVSS
						QAIRDAEKAGPLPHPDMENRG
			}	ł		RVEQRGGYTIEPNNHMRDGVG
						TRWRRAEDVF\PPLIG\VAGHKG
						GR\YKTSV\$VHLAQDLALKGLR
				ŀ		VLLVEGNDPQGTASMYHGWV
						PDLHIHAEDTLLPFYLGEKDDV
						TYAIKPTCWPGLDIIPSCLALHR
		1				IETELMGKFDEGKLPTDPHLML
				ŀ	ł	RLAIETVAHDYDVIVIDSAPNL
		1		1		GIGTINVVCAADVLIVPTPAELF
		1				DYTSALQFFDMLPDLVKNRDL
		l		l		KGVEPDVFIRDKLMERRNRRT
		1		1		GRTEKARIWEVTDRTVRTWIGE
				1		AVAAAAADGVTFSVPVTPHTF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nuclcotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				scquence		
24598	54966	A	24736	1	1400	MDAKMHSGGTAPPGTLSQRQP
24370	34900	ľ^	24730	ľ	1400	SSNSKTNGSKLNPLKDTLLSTPS
			l			CPEGELVNAARRSNGSGNAITL
		1				
		ł				PVWGRDVKFTLEVLRGDSVEK
		l				TSRVWSGNERDQELLTEDALD
1		1			ŀ	DLIPSFLLTGQQTPAFGRRVSGV
		1				IEIADGSRRRKAAALTESDYRV
						LVGELDDEQMAALSRLGNDYR
		l				PTSAYERGQRYASRLQNEFAG
1						NISALADAENISRKIITRCINTAK
						LPKSVVALFSHPGELSARSGDA
						LQKAFTDKEELLKQQASNLHE
						QKKAGCDNLKTCHTSHGSVMA
		ļ				ETAVINHKKRKNSPRIVQSNDL
		1				TEAAYSLSRDQKRMLYLFVDQI
		1		1		RKSDGTLQEHDGICEIHVAKYA
		1				EIFGLTSAEASKDIRQALKSFAG
						KEVVFYRPEEDAGDEKSYEQPS
						KDIRQALKSFAGKEVVFYRPE/
		l	1			VGCRR*KRL*IFSLVYQTCAQSI
						QRALQCTYQPISHSLLYRVTEP
24599	54967	Α	24737	1	1628	MLHITNDQGNASQNHNAMPPY
				1		SCKNGHNLKTVDGGMDAVNG
						EHFYTAGLSQGPLKGSFLLFFNI
		i				LILVCPPIFRVSGNEVFIRDKLM
						ERRNRRTGRTEKARIWEVTDRT
						VRTWIGEAVAPLLLTCDNLKTC
						HTSHGSVMAETAVINHKKRKF
		1				SPRIVQSNDLTEAAYSLSRDQK
				İ		RMLYLFVDQIRKSDGTLQEHD
1						GICEIHVAKYAEIFGLTSAEASK
		l				DIRQALKSFAGKEVVFYRPEED
						AGDEKGYESFPWFIKRAHSPSR
				1		GLYSVHINPYLIPFFIGLQNRFT
						QFRLSETKEITNPYAMRLYESL
						CQYRKPDGSGIVSLKIDWIIERY
						QLPQSYQRMPDFRRRFLQVCV
						NEINSRTPMRLSYIEKKKGRQT
		1		1		THICDNLKTCHTSHGSVMAETA
1		1				VINHKKRKNSPRIVQSNDLTEA
1						AYSLSRDQKRMLYLFVDQIRKS
				1		DGTLQEHDGICEIHVAKYAEIF
1						GLTSAEASKDIRQALKSFAGKE
1				1		VVFYRPE/VGCRR*KRL*IFSLV
						YOTCAOSIORALOCTYOPISHSL
1		l		1		LYRVTEPVYTISAY
		_		L		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24600	54968	Α	24738	1295	1675	SVTSCLKVTSVCLTSGRRFLQIR
		1			1	ISDGTLQEHDGICEIHVAKYAEI
		1		ŀ	İ	FGLTSAEASKDIRQALKSFAGK
						EVVFYRPEEDAGDEKGYESFP
				i		WFIKRAHSPSRGLYSVHINPYLI
		1				PFFIGLQNRFTQFRLSETKEITNP
		ĺ				YAMRLYESLCQYRKPDGSGIVS
		l			ł	LKIAWVIERYQLPQSYQRMPDF
		1		i		RRRFLQVSDFLMAPYRNIDGIC
						EIHVAKYAEIFGLTSAEASKDIR
i		1				QALKSFAGKEVVFYRPEEDAG
						DEKGYESFPWFIKRAHSPSRGL
		l				YSVHINPYLIPFFIGLQNRFTQFR
		1	1		İ	LSETKEITNPYAMRLYESLCQY
		l				RKPDGSGIVSLKIDWIIERYQLP
l						QSYQRMPDFRRRFLQVCVNEIN
ŀ						SRTPMRLSYIEKKKGRQTTHIE
		1				HDGICEIHVAKYAEIFGLTSAEA
						SKDIRQALKSFAGKEVVFYRPE
						EDAGDERGYESFPWFIKRAHSP
						SRGLYSVHINPYLIPFFIGLQNRF
						TQFRLSETKEITNPYAMRLYQS
						LCQYRKPDGSGIVSLKIDWIIER
l		l				YQLPQSYQRMPDFRRRFLQIRK
		l				SDGTLQEHDGICEIHVAKYAEIF
						GLTSAEASKDIRQALKSFAGKE
		1				VVFYRPEEDAGDEEGYESFPWF
	İ			l		IKRAHSPSRGLYSVHINPYLIPFF
		1		l		IGLQNRFTQFRLSETKEITNPYA
				l		MRLYESLCQYRKPDGSCIVSLK
						IDWIIERYQLPPRYQRMPDFRRR
				i		FLRVWVQEINSRTPMPPLLPWE
24601	54969	Α	24739	1	1986	

SEO ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		١.		sequence		
24602	54970	l <sub>A</sub>	24740	1	398	MVDSLIARVGVMARGNAITLP
24002	34970	l^`	24740	<b>'</b>	376	VCGRDVKFTLQVLRGDSVEKT
	i	l		į.		SRVPPGNERDQEMLTEDALDD
		1				LIPSFLLTGQQTPAFGRRVSGVI
	1	1		i .		HTADGSRRRKAAALTEKFCIVD
	ı	1				NPPGEIRKRACLFIPTHALNLNR
		1				FEDICYAHOLPDGGLLSARVGV
		1				MARGNPILSPVCGRNVKFTLKV
		1				LRGDSVENPSRVWSGNERAQE
	ı	l				
	1	l				LLTEHALDDLTLSFLLTGQQTP
	l	l				AFGRRVSGVIEIADGSRRRKAA
						ALTESDYRVLVGELDDEQMAA
	I	l		1		LSRLGNDYRPTSPYERGQRYAS
	1					RLQNEFAGNISALADAENISRKI
						ITRCINTAKLPKSVVALFSHPGE
	1	l				LSARSGNERDQELLTEDALDDL
		l				IPSFLLTGQQTPAFGRRVSGVIH
		l	1			TADGSRRRKAA/DTYRKILYR**
21500	54971	١	24741	1450	3699	PPGGD*KKSVPFYSNTCAQSQP
24603	349/1	A	24/41	1430	3099	RNPLQIPKQQQNEKYQVPQFDQ
		İ				STIKNIESAKGLDVWDSWPLQN
		l				ADGTVAEYNGYHVVFALAGSP
		l				KDADDTSIYMFYQKVGDNSIDS
	i			1		WKNAGRVFKDSDKFDANDPIL
	1					KDQTQEWSGSATFTSDGKIRLF
				1		YTDYSGKHYGKQSLTTAQIFQ MITVFITSLCNTTSSHICLHISWV
		l				TTMHGAMPHPGHIPTAPASLPA
				İ		KVKGSGKVCWLAFPLSELTPPS
			l	1		
						ALQVNVSKSDDTLKINGVEDH KTIFDGDGKTYQNVQHFIDEGN
		1	1	ı		YTCGDNHTLRDPHYVEDKGHK
		1		ļ		YLVFEANTGTENGYQGEESLFN
		1		l		KAYYGGGTNFFRKESQKLQQS
		1				AKKRDAELANGALGIIELNNDY
	1		l	1		TLKKVMKPLITSNTVTDEIERA
		1				NVFKMNGKWYLFTDSRGSKM
	1			ŀ		TIDVPQAKSTNVVITSYMTNRG
				ŀ		FFEPEGSKGSITAAVMAKGLSV
			ľ			ASGNSTPERRRATANENVOPRA
	1			ŀ		IRDAEKAGRLPHPDMEIRGRVE
						ORVGYTIEOINHMRDVFGTRLR
				į		
						RAEDVFPPVIGVAAHKGGVYK
				I		TSVSVHLAQDLALKGLRVLLV
			1	į .		EGNDPQGTASMYHGWVPDLHI
		l	1	l		HAEDTLLPFYLGEKDDVTYAIK
		l		1		PTCWPGLDIIPSCLALHRIETEL
						MRKFDEGNLPTDPHLMLRLAIE
		l				TVAHDYDVIVIDSAPNLGIGTIN
		l		1		VVCAADVLIVPTPAEQLDIS\SS
		L_	l			RQFFD\ILRDLLKNVDLKGFEPD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24604	54972	A	24742	li	3899	MVDSLIAPVGVMARGNAITLPV
		1				RATNVKFTLEELRGDSVEKTSR
		l				VWSEKKDRGGQEAEVSGEGSS
		1		İ		QTKSLQKRQHSNGSDFTARGF
i		l		1	l	HTGQALVRSKGTYWNPTGLGL
				i		SHGTEDPNLIGEELDAGEWDK
		1			i	NTRKELAVLNVRRDFADFGTTI
						KQDFRLLGQTSVDRLLQLSQG
		1				QAVKGNQLLPVSLVKRKTTLA
		l				PNTQTASPRALADSLMQLARQ
		1				VSRLESGQFWVFEGGLRVLVR
						ERVTTPKGT\ASMYQGW\VPD
24605	54973	Α	24743	268	2617	RMFRMKLMETL\SQCINAGHE
						MTKAIAIAQFNDDSPEARKITR
		l				RWRIGEAADLVGVSSQAIRDAE
		l		1		KAGRLPHPDMEISGR\VKHGVG
		1				YTIEQIYL\TRDVFGTRLPRAED\
		l		1		VFPPVIGVAAHKGGVYKTSVSV
		l				HLAQDLALKGLRVLLVEGNDP
		1		ł	1	QGTASMYHGWVPDLHIHAEDT
		l		}		LLPFYLGEKDDVTYAIKPTCWP
		1		1		G\LDIIPSCLALHRIETELMGKFD
		1		ł		EGKLPTDPHLMLRLAIETVAHD
		l			İ	YDVIVIDSAPNLGIGTINVVCAA
		1		1	i	DVLIVPTPAELFDYTSALQFFD
		l			i i	MLRDLLKTVDLKGFEPDVRILL
		l				TKYSNSNGSQSPWMEEQIRDA
		l				WGSMVLKNVVRETDEVGKGQI
		ĺ				RMRTVFEQAIDQRSSTDTSLSTP
		l				AAPMVDSLIARVGVMARGNAI
		1				TLPVCGRDVKFTLEVLRGDSVE
		1				KTSRVWSGNERDQELLTEDAL
		l				DDLIPSFLLTGQQTPAFGRRVSG VIEIADGSRRRKAAALTESDYR
		ı				VLVGELDDEQMAALSRLGNDY
						RPTSAYERGQRYASRLQNEFAG
		ŀ				NISALADAENISRKIITRCINTAK
		į.				LPKSVVALFSHPGELSARSGDA
		1				LQKAFTDKEELLKQQASNLHE
		1				QKKAGVIFEAEEVITLLTSVLKT
			l			SSASRTSLSSRHQFAPGATVLY
		1	l			KGDKMALNLDRSRVPAYIIRSY
						IRCGRKGFASAGVGGCRGWLN
		1	l			YAASEQIVLRVHHMRCEIPHRC
		Ц	<u> </u>	L		THIS EQUALITY THINK CEIF THE

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, =possible nucleotide insertion)
1				sequence		
21/0/	14.004	<u> </u>	2.011		laca.	Lamparin and arrival and arrival
24606	54974	A	24744	1	2624	MTPQKRGYLCKAGARNLDVLL
						EGCEIFLLIPMGMRSESGVSDA
		l		ĺ		DTVSAPRKKHSLEQGIFRAQVE
l	i					ASKPSSVPACIVVGNVSGYQFP
	1	1				PGTSYGPDSESELGYREKFMSF
	1				i	VGTMNEAGNPPFSANYSKAKK
	1					PKPRMFSLIERMSPSPRVGSQSV
Ì	1	l				QSSCFRDIPFEQDQMFPHPSLVL
	i .					RSKEQQPLLTQSIGVNSARKAM
l	1					AKRRRHRRRRRVSGVLPIA\VG
		l				VPS*SVHLPKVLSCSGASWI*QS
	1					CISRLGRLSPTRLMNVA\GNDP
	1				1	QGTASMYHGWVQILHIHAEDT
	1				1	LL/PFY/LGEKDDVTYAIKPTCW
						PGLDIIPSCLALHRIETELMGKF
						DEGKLPTDPHLMLRLAIETVAH
						DYDVIVIDSAPNLGIGTINVVCA
						ADVLIVPTPAELFDYTSALQFFD
1			İ		İ	MLRDLLKNVDLKGFEPDVRILL
	1					TKYSNSNGSQSPWMEEQIRDA
	1				l	WASMV*K\NNVRNANEVGKDT
						LKGFYSLKERPLGNHFLGPNNR
1	1					PFGKIMKHKTPAFGRRVSGVIEI
	ł					ADGSRRRKAAALTESDYRVLV
ł						GELDDEOMAALSRLGNDYRPT
			ĺ			SAYERGORYASRLONEFAGNIS
						ALADAENISRKIITRCINTAKLP
			İ		İ	KSVVALFSHPGELSARSGDALO
					İ	KAFTDKEELLKQQASNLHEQK
	1					KAGGFCRFRNHHOTGFSPAGA
						NORGPLAATLSGPGGEGQSAV
l	1					ARLTGEKKNHPGAQYANRLSP
24607	54975	A	24745	1	385	MWOAISRLLSEOLGEGEIELRN
24007	3.773	ľ	247.45		303	ELPGGEVHAAWHLRYAGHDFF
						VKCDERELLPGFTAEADOLELL
						SRSKTVTVPK\AFILGOOIARLH
	1			1		OWSDHAVLKIVTTGVL*GAYV
						GAPSIWKCPERITLPGYHO
24608	64076		24246		1200	GATSIWACTERITETGTHQ
24608	54976	Α	24746	I	1209	

	n of first   codon for last amino acid   *-Stop codon, /-possible nucleotide
	for peptide of peptide sequence deletion, \=possible nucleotide insertion)
	ce
24609 54977 A 24747 I	857 MHACRSELLGEGEIELRNELPG
	GEVHAAWHLRYAGHDFFVKC
	DERELLPGFTAEADQLELLSRS
	KTVNRALKVWA\VGADRELTV
	FLV\MDYLPPRPLDAHSAFILGQ
1 1 1 1	QIARLHQWSDQPQFGLDFDNA
	LSTTPQPNTWQRRWSTFFAEQR
	IGWQLELAAEKGIAFGNIDAIVE
	HIQQRLASHQPQPSLLHGDLWS
	GNCALGPDGPYIFDPACYWGD
	RECDLAMLPLHTEQPPQIYDGY
	QSVSPLPADFLERQPVYQLYTL
	LNRARLFGGQHLVIAQQSLDRL
24610 54978 A 24748 1	2094
24611 54979 A 24749 1	1209 MVREDLYTYYSCNTSREQQQL
	ENPMESPLCSKEINQGGVFMLA
	GKASDTLLAGGTMNNLGGEDS
	DTIVENGSIYRLGTDGLQLYSS
	GKTQNLSVNVGGRAEVHAGTL
	ENAVIQGGTVILLSPTSADENFV
	VEEDRAPVELTGSVALLDGAS
	MIIGYGAELQQSTITVQQGAPP
	VKLRFALYGRPGFTKCGWMLG
	YKPAHDKTLLRITRGHDFACYQ
	RTTQLARQRGLKVCSHLIVGLP
	GEGQAECLQTLERVVETGVDGI
	KLHPLHIVKGSIMAKAWEAGR
	LNGIELEDYTLTAGEMIRHTPPE
	VIYHRISASARRPTLLAPLWCE
	NRWTGMVELDRYLNEHGVQG
	SALGRPWLPPTENSLMKQIRLL
	AQYYVDLMMKLGLVRFSMLL
	A\WPSSFLPLWYKWR
24612 54980 A 24750 200	310
24613 54981 A 24751 I	4398 MCIPVPPIATGDRHFIAVAHIHQ
	NSRRSPVDSDESSTRSAVAAHP
1 1 1 1	ATRDFNVRALYTVTSSVSPTKP
1 1 1 1	PLPATFRQGRYRPAEPCVGSIG
	VIPQSSCVLANGSTGAQSAFGDI
	IRAGPPSAGLVPVAQATVIETFL
	RYQTINERDFWAARNRLLKQP
	YGSRILRFTEAYAIAERAAQSV
	LRISLADAFNVDGPRPAEDSIFR
	LLRMPGSAVQRPDRRHNRRCQ
	YRPAMAFPAGSKANKQRQYNG
	GRQFYRKRICWL

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24614	54000	<u> </u>	0.4750	lose	10160	Indiana in the control of the contro
24614	54982	Α	24752	939	2169	RPASPPARDGARWSTGLHSSYS
			l	I		AMRCRNASTCAEDSQLS/DPLA
1	1	1	l	1		R/PSVASGPYRVTSWKLGQNIV
1	1			l		YCRVKDYWPATLTVNRGRWN
	1	1	1	I		FDTIRSDYYLDDNVAFEAFKAG
1		1	l	l		AFDLRMENDAK\NWATRYT/G
l			l	l		DRRVREAITLAFDFEWMNKAL
	1		l	1		FYNAWSRTNSYFQNTEYAARN
	1	1	l	1		SPPPPEL\VLLPPIKKVLPSEVFP\
		1	l	[		QIYQPPVSKGDGYDRDNLLKA
		1	l	1		DKLLNEAGWVLKGOORVNAT
1	1		l			TGOPLSFELLLPASSNSOWVLPF
1			l	1		OHSLORLGINMDIRKVDNSOIT
1	1		1	l		NRMRSRDYDMMPRVWRAMP
i			l	1	1	WPSSDLQISWSSEYINSTYNAP
1			l	1		GVOSPVIDSLINOIIAAOGNKEK
1			l	1		FLPLGRALDRVLTWNYYMLPM
1			l	1		WYMAEDRLAWWDKFSOPAVR
		1	l	I	l	PIYSLGIDTWWYDVNKTAKLPS
		1	l	l		
24615	54983	l.	24753	12	450	ASKQGE
24615 24616	54983	A	24753	2	203	DI DDA ADVI BROCOVE A BUTTO
24016	24984	^\	24/34	ľ	203	PLPDAARYLRRQCQKPARVHDI
1			1	İ		FAAMHTAHRSRPDRPSLPDDAF
1			ĺ	I		SSLRL*FFQPEHYPRQYQHRHL
1			l	I		CS*ERYQHQALSRTRFRSG*PTP
1			l	I		SVPETSTGSRYFCGNAHSAPKP
1		1	l	I		PRPPITTGRRVFFTTPLILPTRAL
		L.			1212	PASISTPASL
24617	54985	A	24755	1016	1219	
24618	54986	Α	24756	1695	2021	PDTFSMLPATRFLPPLQFPDLQH
1		1	l	I		PEAAQYAHWLPASGCFLFLPPP
l		1	l	I		GSGTVFLLPYPSPRLCRSTSETP
1	1	1	l	i		LPAITR*HFPVTDRVQK*ITRRR
		_				KPRHKPFYPDDLQRDAR
24619	54987	A	24757	1671	1911	
24620	54988	Α	24758	513	1104	CWRNCARCERNLVFRCQYGIY
I	1		İ	1		WCDNAANRSCRTKRDQPRLVE
l	l	1	Ì	1		W*KSACSHHR*LH\PCGHCRQF
			1	1		MNELNSGLDLRIHLPGREAHAL
l		1	1	1		RDYLPDAFGPKDLEIKTLLMDE
			l	t		QDHGYALTGDALSQAAIAAAN
			Ì	1		RSHMPYSKSPSGVALECKDGRI\
l				I		LVAATLKTPH/INPTQICGQPWS
	I	1	l	l		MHRRALTSKFGAHGDRRTSAG
			1			FRRN
2462I	54989	A	24759	131	759	
24622	54990	A	24760	662	889	LPIKCRAARGVPARCLRWTRW
	1	1	l	1	[	ALRRILPPLRNRSRAASRWRAS
			1	1		PGARK*WMPSLQAVWAAPMR
	1		1	1		VTRLPAWLRWKC
	L		L	L		

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24623	54991	I <sub>A</sub>	24761	1	1518	MQRRDGDIALIYRSKIGPRTSIT
	.,,,,	1		ľ		FTARRANPVQRIAARILLRDHLI
					1	CRMSCTAARHANTFDLIQGSG
		1				ADNFAAIYGGANDRITRADAA
					1	VAGAGNWHRFGISNGKLSPRSI
	ŀ					TRTSSPARAMTKMPYNPOASG
		1				GSNLAVSNPELDKLIAARVAAL
						RAANPDASASVPVELVTASASY
		1				PRSLMRCNRILSPGSARGQMIIT
						ALFIYAGNPIACVAALEVLKVF
		l				EQENLLQKANDLGQKLKDGLL
1						AISFWRAKWLKIFVVCGAEIFV
						EYVLKNKVIWLLCFANIFLYVV
						RIGIDQWSTVYAFQELKLSKAV
						AIQGFTLFEAGALVGTLLWGW
						LSDLANGRRGLVACIALALIIAT
						LVIPFHMVFHPNEQYAYCVNEL
						NSSVDVWELKDPHGNIECVQT
						LDMMPENFSDTRWAADIHITPD
						GR\HLYACDRTASLITVFSVSED
				1		GSVLSKEGFQPTETQPRGFNVD
		1				HSGKYLIAAGQKSHHISVYEIV
		1				GEQGLLHEKGRYAVGQGP\MW
24624	54992	Α	24762	3	486	ACRRALKD\AGVEADEVLEVV
	-					MVGGSTRVPLVRERVGEFFGRF
	ł	Į				PLTSIDPDKVVAIGAAIQADILV
		1				GNKPDSEMLLLDVIPLSLGLET
				1	ŀ	MGGLVEKVIPRNTTIPVARAOD
						FTTFKDGQTAMSAGGAHIRVTF
				1		OVDADGLLSVTAMEKSTGVEA
						SIOVKPSYG
24625	54993	В	24763	1	2469	5127111515
24626	54994	A	24764	1255	1761	ESFOVRTHGTGEPVNKTRTSOA
						VSHHNOSCOVDOCVPCAVIAG
						NVFPRHNVQGQH*RNCQQTNR
	1	l	1			SSINDFTKLSNTRPISVHSLGIKO
		1	1	I		SRPDRHAKASPVVTAVKACEO
		1				VIAAVNQIADDHQPDRTKQRN
						AIVVADDLPDLFPVHFFGVDHQ
		1				QHNHRDKKQPGEDLLCC
24627	54995	A	24765	1	3141	QUINTERDARQI GEDELECC
24628	54996	B	24766	1	3969	
24028	24990	l <sub>B</sub>	Z4/00	11	3707	

SEQ ID NO:	SEQ ID NO: of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
24629	54997		24767	643	2082	GAHAMDKHESTTPAEKNLEVI- LALIGIWYNNFFDKNDALVMV DDFPRRSGLSWKIVVVPHEYW VGGSTRVPLVRERVGEFFGRPY VGGSTRVPLVRERVGEFFGRPY UTSIDPDKVVAIGAAIQADILVG NKPDSEMLLLDVIPLSLGLETM GGLVEK VIPRNTTIPVARAQDF TTFKDGQTAMSIHVMQGEREL VQDCRSLARFAICVVFRRYRLA GAHIBVTFGVDADGLLSVTM EKSTGVEASIQVKPSYGLTDSEI ASMIKDSMSYAEQDVKARMLA EKSTGVEASIQVKPSYGLTDSEI ASMIKDSMSYAEQDVKARMLA EKSTGVEASIQVKPSYGLTDSEI ACMIKOTTOTAGTDHRA DVTEVASDFRVQTTNADSALF HVRSAQQHIHQLLHFTTHLLCQ LAGLIDHVVFQQIATNFADQVG MVGAFTTGKDLCHFHGGFTHA EELHKAGVEAGKVTGQAKVEG MRVQAFVLQONGADHLRTFW HLNTHRVFHRRGVGGAVGKA ADPAHTVREEGHFVHTIFTGFQ FFHAAMDVEV
24630	54998	Α	24768	1	502	

SEQ ID	SEQ ID NO:					Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	or pepude sequence	desentin, (-possible nucleotide insertion)
24631	54999	IA	24769	li	2345	MFMIGALVLALVMTLITFMVL
		1				RRIVIRPLOHAAORIEKIASGDI
		1	Ì			TMNDEPAGRNEIGRLSRHLOG
	i	1			1	MQHSLGMTVGTVRQGAEEIY
	i					GTSEISAGNADLSSRTEEQAAA
	l	1			1	EQTAASMEQLTATVKQNADN
						HHASKLAQEASIKASEGGQTV
		1				GVVKTMGAMSTSSKKISEITG
		1				HOOGIRTPVTAVKGRCPGPLE
ı						GDTDTAYYALRDIFALTPNRK
				1	i	LAIPVSTDNPGKPCKDPSLSKI
					VSNRVPQRVPCYDSPRSSSVN	
						NRLPGKDFHDDERSMGRSGS
	ľ				VLALAYPPKIALSDADSVTVN	
	1			l	AGWQGEISREQFNELIAPLVK	
	1				TLLACRRALKD\AGVEADEVL	
	1				VVMVGGSTRVPLVRERVGEF	
				i	GRPPLTSIDPDKVVAIGAAIQA	
						ILVGNKPDSEMLLLDVIPLSLO
					ETMGGLVEKVIPRNTTIPVAR	
					ODFTTFKDGQTAMSIHVMQG	
						RELVODCRSLARFACVVFRRY
					l	LAGPVDLFLRPWEVDISRRTS
						DSPLPVQVLEASPKGHYTQLV
				i		QPLGWYNEPLTVVMHGDDAI
		l				ORGERLFVGLQHARLYNGDE
						ASRDPGFLMARFVTCRPDKTF
				1		LRIRQHHVWIEIVSTLEQTIGN
				1		PLVKLORMGPDNGSEVWLKL
						GNNPAGSVKDRAALSMIVEA
				1		KRGEIKPGDVLIEATSGNTGIA
						AMIAALKSYRMKLLMPDNMS
24632	55000	A	24770	313	676	DCOPRSSAFTKPDATTAGHYL
24032	33000	1.	21770	3.5	10,0	GSGGRMTQLAIGKPASLGAH
				ì		REGGENFTFFSLNADGEKL/C
				1		DANGO/EHRYDLPGHSGDIWI
						G/YLPDARPALRMVSRH/GP/W
						PAGGMLTPAVVMILPRK
24633	55001	A	24771	1760	2053	FRVHGKNAFSH*IKPAAMLLP
24033	33001	Ι.		1.700	2005	SPVTRSVKPTQCGSAPSKRPN
						TLLPSITLMPPLPVTDLKLGCI
						PCSSDCFSFITFSISCATRLSVN
			1			LPAAGO
24634	55002	A	24772	296	732	
24635	55003	Α	24773	606	695	
24636	55004	Α	24774	1520	1737	PVHGLCGTRRDAGS\PAIYPFF
		1	1			LML/RFWLR/TCTVPGWPLDD
		1	1			SLAGEAFSAFTTDAAVWSLIS.
		1	1			TAMLAVQAG
24637	55005	A	24775	1	1049	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleutide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24638	55006	Α	24776	248	933	RTTMRAVSRPKYCSRERWL/HY
						DFTVARFDENASCGTFAATSAV
		l				VLIFSHCLRLLCRVVVLVTRVN
		1				FQFTEHSTTQRAFWQHAFNRDF
		1				NHTLRTASNHLFKGRLFDTTDV
				*		AGVVIVHFVSTLVAGYSNFVSV
		Į.		1		QNDDVITGIYVRSVFRFVLTAQ
						ATSQFSSQTAQSFTGRVNNIPV
ŀ				l		AFYGFWFSCAKYYRHGARWCS
l		1		i .		NGRKIDQRHTHCFFCPCIRTDFA
		<u> </u>				VEIGKEEFIT
24639	55007	Α	24777	219	3441	DCQPRSSAFTKPDATTAGHYLA
						GHSGDIWHGYLPDARPGLRYG YRVHGPWOPAEGHRFNPAK\LL
						IDPCARQIDGEFKDNPLLHAGH
						NEPDYRDNAAIAPKCVVVVDH
						YDWEDDAPPRTPWGSTIIYEAH
		1				VKGLTYLHPEIPVEIRGTYKAL
	1					GHPVMINYLKOLGITALELLPV
						AQFASEPRLQRMGLSNYWGYN
						PVAMFALHPAYACSPETALDEF
					1	RDAIKALHKAGIEVILDIVLNHS
	1	1				AELDLDGPLFSLRGI
24640	55008	Α	24778	454	586	
24641	55009	В	24779	130	1557	
24642	55010	В	24780	92	2808	
24643	55011	A	24781	2	1051	WRLRPITPLWRWQKIFFPDA/R
				l		YGIRFRNLETGNWYPELLDNVE
		1			1	PSFVWANDSWIFYYVRKHPIFI
		1		i	1	DLTTEELAASSYHESWVLEDRII
		1		l	i	KDGSYNIDQGVGVRAISGEKTG
		ı	İ	l	I	FAYADQISLLALEQSAQAARTI
	į.	1				VRDSGDGKVQTLGAVEHSPLY TSGRSGWQSMSREEKLDILRRV
		1			1	DKVAREADKRVQEVTASLSGV
		1		l	1	YELILVAATDGTLAADVRPLGK
		1				VLLVGQSPNAELSARAKQIAM
			1	I	1	GVDGANEVYNEIRQGQPIGLGE
1				I	1	ASNDTWITTKVRSOLLTSDLVK
		1	l	l		SSNVKVTTENGEVFLMGLVTE
1			1			REAKAAADIASRVSGVKRVNR
				l	1	TLFLGYAVMFSELILAPVTPSNS
24644	55012	В	24782	1	2079	
	1-7	1				L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /-possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24645	55013	A	24783	1	3214	MGAEKQLIPDGCGVSYIPSSGP
						LDKWRALLFIQPDYRADPVYR
		1				QHTITDLRILYDYTENFVLPLSH
	İ					DEVVHGKKSILDRMPGDAWOK
		l				FANLRAYYGWMWAFPGKKLL
						FMGNEFAQGREWNHDASLDW
						HLLEGGDNWHHGVQRLVRDL
				1		NLTYRHHKAMHELDFDPYGFE
						WLVVDDKERSVLIFVRRDKEG
1	j					NEIIVASNFTPVPRHDYRFGINQ
						PGKWREILNTDSMHYHGSNAG
		l	1			NGGTVHSDEIASHGRQHSLSL
24646	55014	١.	24704	1	1079	
24646	55014	Α	24784	11	1079	MKQIKELQRLLGKKTWKMNSS
		i		1	\	KKPLNMDGPKVDSARALIGRG
				1		WGTHKFYRCDCIKPTIKTNTSS
				i		RGNEKNPDGSHNNIHNSWFRI.
						LTSQNMCKVPESPAILVDVGQS
1						AETVQSVLTVKVEHPILRIAVA
					l .	GYACQRQSCLLAMRHPQYAIN
1						SSKVVKPCSTARQPFQKAYAYP
						PVVRAFPAINVGISGNKLTNDR
						RYGRTSNKAWGSGDGHTLSEC
						GWAPLAAGDNVGKVSYQTLG
						YYQRILGYIFYPIAW/VMGVPSS
		1		ŀ		EALQVGSIMA/TKLVSNEFVAM
	1				1	MDLQK/IASTLSPRAEGIISVFL/
				i		VSFANFSSIGIIAAAAECRDWFG
	1	1				ILAAQTLAVRAALRVVQAERD
						CVLSMEQHQCARGRQR
24647	55015	Α	24785	121	258	LTEPRHYCRSIQFFGLTNMGQN
						CYQLS*E*FIGTLNHIRRIPQPLS
24648	55016	Α	24786	1	470	WGSGDGHTLSECGWAPLAAG
l						DNVGKVSYQTLGYYQRILGYIF
	1					YPIAW/VMGVPSSEALQVGSIM
1				1		A/TKLVSNEFVAMMDLQK/IAS
	1					TLSPRAEGIISVFL/VSFANFSSIG
ì	İ					IIAAAAECRDWFGILAAQTLAV
				l		RAALRVVQAERDCVLSMEQHQ
						CARGRQR
24649	55017	Α	24787	1324	1539	GWCGCEVPFKITDGLACLLQSP
				1		QGHQTFCFT*HNLGLQPFWIPM
l	1			l	1	CLCSVNVFKTRLILTQVEEGCR
1		1	1		1	AVSMTQ
24650	55018	A	24788	71	403	RTTFIFDLSSGWPFILWQVLSPN
				1		*RCLQVGLFAQIPTGRMGCLHR
l	1	1				FLTCRMGVCTDSSPAGWAVTLI
				1		RMSPIFPASSKSTLSAALLNTSC
1	1	1				LHSRDLEPPGAFRSGGPVLL

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24651	55019	A	24789	309	1653	SVLMDESDIYRNNRERFFCVFL
2.051	330.7	1	24707	507	1,055	LVGVPARGACLGGGGGGCGLCV
						RYLRSAGWGCGRRFPCVGDN
		1		i		WSAILAHIGKPEELDTSARNAG
						ALTRREIRDAATLLRLGLAYG
						PGGMSLREVTAWAQLHDVATL
	l .	1				SDVALLKRLPKCRHWFGILAG
					1	QTLAHIRVLPVIIRAIGFLLSKV
		1			i	NGMGKLESFNAVSSLILGOSEN
						FIAYKDILGKISRNRMYTMAAT
				1		AMSTVSMSIVGAYMTMLEPKY
						VVAALVLNMFSTFIVLSLINPYR
				1		VDASEENIQMSNLFEGQSFFEM
						LGEYILAGFKVAIIVAAMLIGFI
		1				ALIAALNALFATVTGWFGYSIS
	1	1				FOGILGYIFYPIAWVMGVPSSE
	1	1				ALOVGSIMATKLVSNEFVAMM
	ŀ	1				DLQKIASTLSPCAEGIISVLLDSF
	l .					ANFSSIGIIARAVKGLNEEOGNE
						GSRFGLKLVYRSTLVS\VLSASI
			l			AALVLSDHT
24652	55020	Α	24790	754	957	
24653	55021	В	24791	75	3707	
24654	55022	В	24792	75	264	
24655	55023	В	24793	344	461	
24656	55024	Α	24794	1	2203	
24657	55025	Α	24795	20	198	
24658	55026	Α	24796	1239	1409	RGPPTPFFHVWEWQQRPLSVW
	1	1				EADHV*RRDEGACPRMVARAR
						NCRPKLLELRAESQ
24659	55027	Α	24797	1325	1564	
24660	55028	Α	24798	323	694	TGYARQKPVPALGWWTNVSAS
	ŀ					TAPPIILPSITGSRLAVIKWLTVS
	1	1				AGKLSALLCASCQKGGGAKWR
	1	1	1			MAMANGITNKLAIQCSPPHSTI
	i	l				EKFTSQSQHVLA*GVSAVRSM
		l				VSSG*GRRPLPALG
24661	55029	Α	24799	4708	4878	
24662	55030	A	24800	20	199	
24663	55031	A	24801	743	1186	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
24664	55032	A	24802	1394	2651	RLMTCRHEGSLGDDPGINGQL WDVNRIDVTAHEGTWERWTV ARDEPQAFHIEGVMFQIRNVNG AMPPPDBGWKDTVWVDGQV LLVYFGQSWAHPPFYNGVO LLSWPQYWAACEGPAPFLDMS REEMDQLGWDSCDIILVTGDA YVDHPSFGMAICGRMLEAQGF RVGIIAQPDWSSKDDPMRLGKP NLFGVTAGNMDSMINSYTAD RRLRHDDAYTPDNVAGKRPDR ATLVYTQRCKEAWKDVPVILG GIEASILRRTAHYDYWSDTVRRS VLVDSKADMLMFGNGERFLVE VLVDSKADMLMFGNGERFLVE VLVDSKADMLMFGNGERFLVE VLVDSKADMLPGVSTRLDTTPGKI DPIPHPYGEDLPCADNKPVAPK KQEAKAVTVQPPRREVWEKTY VLLPSFEKVKGDKVLVAHASRI LHHEINPAVPAH

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	tocation of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24665	55033	A	24803	11	2126	MOEKQVLAAYVQPSSEGOSRS
				ľ		RRARRVSOCHOHSAAAASLAR
		1		1	1	TAASATADSAHPPCPRSAARAG
		1				ATMHVKKYLLKGLHRLOKGPO
						YTYKELLVWYCDNTNTHGPKR
		1				IICEGPKKKAMWFLLTLLFAAL
		1				VCWOWGIFIRTYLSWEVSVSLS
	1	1				VGFKTMDFPAVTICNASPFKYS
	1					KIKHLLKDLDELMEAVLERILA
		1		l		PELSHANATRNLSFSIVNPTP\L
	ŀ				ŀ	
	ł					VLIDERNPHHPMVLDLFGDNH
		1		ł		NGLTSSSASEKICNAHGCKMA
	ĺ				l	MRLCSLNRTQCTFRNFTSATQA
				l		LTEWYILQATNIFAQVPQQELV
	1					EMSYPGEQMILACLFGAEPCNY
	l					RNFTSIFYPHYGNCYIFNWGMT
						EKALPSANPGTEFGLKLILDIGQ
			ļ			EDYVPFLASTAGVRLMLHEQR
	1	l			l	SYPFIRDEGIYPMSGTETSIGVL
					ŀ	VDKLQRMGEPYSPCTVNGSEV
					ł	PVQNFYSDYNTTYSIQACLRSC
		1		l		FQDHMIRNCNCGHYLYPLPRG
		1				EKYCNNRDFPDWAHCYSDLQ
	1	ľ				MSVAQRETCIGMCKESCNDTQ
					1	YKMTISMADWPSEASEDWIFH
		1				VLSQERDQSTNITLSRKGIVKLN
						IFFQEFNYRTIEESAANNIVWLL
		1			İ	SNLGGQFGFWMGGSVLCLIEFG
i		1				EIIIDFVWITIIKLVALAKSLRQR
						RAQASYAGPPPTVAELVEAHT
		1				NFGFQP\DTAPRSPNIGP\YPGEQ
ĺ						ALPNPG/TPAPPNYDSWRLQPL
24666	55034	Α	24804	905	1051	
24667	55035	A	24805	1	1257	
24668	55036	Α	24806	1	3116	MPIITLPDGSQRSFDHPVSVAEV
1	1					aQSIGAGLAKATLAGKVDGRLI
	ì	1				ITPKDEEGLEIIRHSCAHLVGHA
l		1				VKQLYPTAKMVIGPVIEEGFYY
	İ					DIFFERPFTPEDMAAIQQRMRE
ł						LIDKDYDVIKKMTPRAEVIELF
		1				KLQAAPDRRHAGREGHGLYFH
		1				EEYVDMCRGPHVPNTRFLKAF
		1				QLTKISGAYWRGDSKNEQLQRI
		l	l	l		YGTAWADKKOLAAYIQRIEEA
		ı		l		EKRDHRRIGKQLDLFHLQEEAP
1	1	1	l	I		GMVFWHPNGWSVYO
24669	55037	A	24807	1924	2790	
24670	55038	A	24808	1	1185	
	1	1		1	1	<u></u>

	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24671	55039	Α	24809	823	5990	MARPVPLWSSWGLRTSYLPKP
						LSVNPSRNDGSA/FAGTHPLGV
						TRGTLAGPAFVRLSLVFLLRRV
						LAGSSLLPLLMG*QK/PEVLLPD
						GASEHDYSLRPSDVKRLQNAD
						LVVWVGPEMEAFMQETGKQIT
						RSEVSPLCRRVAFNKFNRDPQP
						RYDWGRLTVLYSGSSLSGGRY
						VVIIRECHCQLSKPLRLEQNEIR
						SLLPEASEPIDQAAQEDEAIPQD
				ŀ		ELDDKIAGEAGVHEYVVSTGD
						TLSSILNQYGIDMGD
	55040	Α	24810	2305	2550	
24673	55041	A	24811	665	844	ISFNNEVISYRLSGKNQ*LNLG*
				1		EIWLESGAFKKALSPQKHPGLE
1						PSVCTPTPTLVPDR
24674	55042	Α	24812	1	2031	
24675	55043	Α	24813	106	912	HFETVIDQRDFHPVFSHKGIRAT
						PEGNEEKGQRVADVTEQCGRC
ĺ						RCQPPAATLTQAVKNVHVEDL
						PQRVGHETGGGDTGDKQVEVG
						ECAESHTLDALGLRCPEPVMM
						VRKTVRNMQPGETLLIIADDPA
			ļ			TTRDIPGFCTFMEHELVAKETD
l		1				GLPYRYLIRKGG**GLIGFDAAF
		l		l		SPHPNPLPRGARGPIALNVAIGL
		l		l		PYLLRNNLNAFAVTSTVAPVSA
		1		l		STASHNPVIQRGDEEDPFQPQR
			1	l		NSDVLADIGVSGACQFNHLHQ
						AAQVVMG

SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
	of peptide	hođ	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24676	55044	Λ	24814	1	1495	MOKLTLTYAETRRLYGKEVYH
24070	33044	^	24614	l'	1493	RPSRFIGELPEECVEEVRLRATV
				ŀ		SRPVSHORMGTPMVENDSGYK
					İ	
					1	LGQRVRHAKFGEGTIVNMEGS GIAALCHCRVLIATAALMDPDD
						AVELSVTTMMFALCNTRNPRV
						AWDTAPCVPATTAASMAKRAP
	ĺ			l		DMSQSAAPEAGSQKPPRLSCGE
						SLHPLPATKRAYNYCRPVGRIR
						HLCRIRQSMPDATLSRLIRPTTV
					İ	ADPPDKALAPHPAVFTRGDSKF
			i		i	LSHKPARSPAIRYLLASVWGPV
			ļ			GMMAMMLAMTGFVVATRKLR
					İ	RRFGFKSCKNGVLYRISSSSITT
						SPASPIATCRPVLPCWMRMDPM
		i		l		LDMMGMQMLMEKYGDQAMA
				ĺ		GMDHSQMMGHMFLGHKVIRA
		ŀ				NHVGDWGTQFGMLFRIAVETF
		ŀ				KEDWEAVWDQLDDLNLEGKIV
						A*GRDPAAFHIDNGAFTKLSVA
						YALAELVAAVVLDHRRTHPLM
						ADRAAYRGAQAHLFHTLFRQL
						ADKARRAMVNLFTIOTAGFRV
						GQRQFLH
24677	55045	Α	24815	3	707	GCTLGPVKSFSLRVAWLSWPQ
		ĺ				YFYIIERACSPCCGPLAPGEAPH
		l				VNLFHSTHSENKKQPHNSPYHN
		ĺ				DIQPTLNYLPTNQKKQTQQEQS
		l				HNITKQGSQLHDVTQVPYQAY
		l				RSAIVVGLIRRDSVASGIDCRM
		İ		l		RRERLIRPTGRASDNTEFTGTO
		İ		l		NMONITHSWFVOVMIKATTDA
		İ				WLKGWA*AQRRQPDATPG*RR
		İ				YRTISRQFPATSVLYPAQPAHA
						FTGKYTVYCHRLGQILP
24678	55046	A	24816	2	1340	
	55047	A	24817	1	2457	

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown.
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24680	55048	Α	24818	41	1053	QRNQQAHLSF/QHSRDIYDNQV
						HLVRNCNLEMPDQEFIVLYGPS
						DCGKSTTLRMIAGLEEISGGDL
						LIDGKRMNDVPAKARNIAMVF
				l		QNYALYPHMTVYDNMAFGLK
ĺ						MQKIAKEVIDERVNWAAQILG
						LREYLKRKPGALSGGQRQRVA
						LGRAIVREAGVFLMDEPLSNLD
						AKLRVQMRAEISKLHQKLNTT
				l		MIYVTHDQTEAMTMATRIVIM
						KDGIVQQVGAPKTVYNQPANM
						FVSGFIGSPAMNFIRGTIDGDKF
						VTETLKLTIPEEKLAVLKTQEV
	l					CPDHTLDALGLRCPEPV/HDGA
						QNRAQYAAWRNVADYRRRSG
	1					HYPRYSWVLYLYGTRTGC
24681	55049	Α	24819	3	131	
24682	55050	Α	24820	247	492	
24683	55051	Α	24821	2	241	YFFQKSSKTVIGGRRLCSVFRY
						QFSIVLATC*GTFFPSR*STILS*
1						VTQSATSSGREGRSSSRSLNGRF
						MFSIASMMSW
24684	55052	Α	24822	1	1143	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24695	55053	_	24823	242	2151	FSROHVVKMKHCCKNVVILMP
24685	33033	A	24823	242	2131	EPVAEPALNGLRUNLRIVSIVM
				1	1	
	ŀ	1				NFASYLTIGLPLAVLPGYVHDV
		1				MGFSAFWAGLVISLQYFATLLS
		1				RPHAGRYADSLGPKKIVVFGLC
		İ				GCFLSGLGYLTAGLTAVCLSSA
		1		į		VILCRGVQLHPVSDLRIPRGEK
		1		j		AGRNRSVDGCCEKAMSVIIVG
						GGMAGATLALAISRLSHGALPV
		l				HLIEATAPESHAHPGFDGRAIAI
		l				AAGTCQQLARIGVWQSLADCA
						TAITTVHVVELHNVGQRLFA\L
		İ		1		LRKAPGVTLHCPDRVANVART
						QSHVEVTLESGETLTGRVLVAA
						DGTHSALATACGVDWQQEPYE
		ļ				QLAVIANVATSVAHEGRAFERF
						TQHGPLAMLPMSDGRCSLVWC
ĺ		ı	i		1	HPLERREEVLSWSDEKFCRELQ
		ŀ				SAFGWRLGKITHAGKRSAYPL
		ŀ				ALTHAARSITHRTVLIQSERLNE
						KALEKRTLTVWDNSFLIYRQSD
						VSEIRLASLFASFSIRKSKQQAR
						KHTNRRSKDKEKRTROKRKTT
		1	İ			SRORRRRORMVRVGDEQIAGC
						HCLYSTGSPNRQLREAGIELSDI
						VRGNEKARERMKAQYSIAGMT
						SGVVVGTDHAAEAITGFFTKYO
		1				DGGTDINPLYRLNKROGKOLL
		ŀ	1			AALACPEHLYKKAPTADLEDD
24686	55054	A	24824	2	91	THIS TO SHE THE RESTREET
24687	55055	В	24825	1	2346	
24688	55056	Ā	24826	I	1713	
24689	55057	A	24827	i	918	
24690	55058	A	24828	415	816	
24691	55059	A	24829	72	296	FCSATFGRPSDOYONGE*LRWF
_,,,,				l -		GLGSFACOSPAHENPOWKFVL
		l		1		ARRGFARGTAEAHPSQYRHWC
		1	1			KIHRLNPGLR
24692	55060	A	24830	1	620	
24072	22000	1,	2-7050	<u></u>	1	L

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24693	55061	A	24831	1625	2460	GVVNPRQGRVLSMLPAMLLYL
		1				LFFLIOTSLKSNGGKGKLDPTL
İ		l				WMWTVNLIYLALAIVLNLWDT
		l				VPVRRLRASFSRKGARIPFVGC
		l				RETCELPQIELVKMMLGRELDT
						HALORAGRTLLSDKPVAAFKN
		ŀ				YGKKGTIAPFDLEVRPGEIVGL
						AGLLGSGRTETAEVIFGIKPADS
						GTALIKGKPONLRSPHOASVLG
						IGFCPEDRKTDGIIAAASVRENII
	1					LALQAQRG\CYVPFPAKNSKRL
]	İ					PNALSASLAFAHLQLNNRLNFS
						PAAISKKCCFYVGY
24694	55062	A	24832	1	2034	
24695	55063	Α	24833	3839	3970	
24696	55064	В	24834	694	765	
24697	55065	Α	24835	1778	1982	Many I and Doc Lavior
24698	55066	Α	24836	1	268	MTCRLLTRLPGQAGVGKRVSG
		l				VPDELSPLFSVGSRYLQGGVGL
		l				VSPTSGANVGLPVISTNNLVKE
		ļ		510	210	CR\DGKWKMTEKGCLNMFHFK
24699	55067 55068	A	24837	217	713	A PTROPOUNTABRO A MOCKANA
24700	22068	Α	24838	217	1910	AFTFQESVMAFSQAVSGLNAA ATNLDVIGNNIANSATYGFKSG
l		l				TASFADMFAGSKVGLGVKVAG
		1				ITODFTDGTTTNTGRGLDVAIS
		1				ONGFFRLVDSNGSVFYSRNGOF
		l				KLDENRNLVNMQGLQLTGYPA
						TGTPPTIQQGANPTNISIPNTLM
1		1				AAKTTTTASMOINLNSSDPLPT
						VTPFSASNADSYNKKGSVTVFD
1		1				SOGNAHDMSVYFVKTGDNNW
		l				QVYTQDSSDPNSIAKTATTLEF
1	1					NANGTLVDGAM/ANNIATGAIN
	1	l				/GLSFLNSMOONTGANNIVATT
1		l				QNGYKPGDLVSYQINDDGTVV
		1				GNYSNEQTQLLGQIVLANFAN
1		l				NEGLASEGDNV\WSATOSSGVA
						LLGTAGTGNFGTLTN/GSIESGI
		1				AFLVNVPLNPKLKRVNVAGAR
						WHPYGWPAFAERACLIEPEGD
						RRTHHRPNNQRTYSRKTQRQG
		1				GHKTRQDNWEQKGDGGPRPEP
1		1				TDPHOHDANRPTVATKKRRKS
	1	l			1	TOSORNHRNPKGTHPTEEHGT
						AGTTKPPDTPOTDNWTHKGSL
		1		1		REGKANRKRNPLRGAEVAGGG
						KORAKAEROTORGERREDVDE
		1				GEEDRSKKTEK
24701	55069	A	24839	500	653	OLEDIOR IER
24/01	122002	<u> </u>	64037	1200	000	1

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /-possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24702	55070	Α	24840	445	849	GSPGTAGTEVAVVLKIGGVHD
		l		1		KVIGHYRCSTTNTGRVLDVAI/S
	i i	l		ł	1	QNGFFRLVDSNGSVFY/SRNGQ
	1	l		1		FKLDENRNPATGTPPT/IQQGAN
	1	1		1		PTNISIPNTLMQRKPTAGQIVLA
	1	l		1		NFANHEGLASEATTSGLRRNLL
	1	l	1	ł		AWRCM
24703	55071	Α	2484 I	I	453	MKQMDTRGKVINCAFAVSAAF
						ALGDHLGFAAANMNAMIFPMI
	1	İ		ı		VGKLIGGVTAIGVAMMLVPRY
	1					EFIKREISWQSPVFFTPVDKQGG
		ļ				LKEAELKTLILEQYHAAGIEPES
		i		1		VDSGAIIITGESAKTRNAR*TGR
		l				FKRSGTENL\KLEQYHAAGIEPE
		l				SVDSGAIIITGESAKTRNARPTV
ł				1		MALSQSLGDFVVASAGPHLE
24704	55072	Α	24842	370	670	YSSNIMLRVLSRKALILVPSSSP
	1	l		i		VKARKPAMLARRGWRSLSRW
	1	l	İ			AILSLPGPGR*TGRFKRSGTENL
		l				\KLEQYHAAGIEPESVDSGAIIIT
	1	ļ			-	GESAKTRNARPAGMALSOSLG
		1				DFVVARAGPHLESVNAGYRTRI
		ŀ				RNHGHSWFPGEIGIRSQFPHTLR
				1		TGSINGKA WGA
24705	55073	Α	24843	I	1623	
24706	55074	В	24844	1	4986	
24707	55075	В	24845	65	726	
24708	55076	Α	24846	4107	4460	FMSTCGYGGGLSRFPARCLSYQ
		1		1		PLRSWRETVAASG*RRTGSARA
		1	Į.	1		NCFCFSGLTRDFSIWRPETK*SA
				1	1	QTFVVGTWRCGGMGR*IAAVL
		1		1		SRYSTVESGVSSTHHRLPLQPDI
		_				PSGRRL
24709	55077	A	24847	624	895	PLVPAYKQSQRQKPHDYLNRC
		1		1		RKSL*QNSTTLHAKNSQ*IRY*
	1	1	İ	1		WD/SISE**ELSMTNPQPISY*MG
ļ.	1			1	1	KNWKHSL*KLAQDRDALSHRS
						YST
24710	55078	Α	24848	1434	1594	
24711	55079	Α	24849	1294	1488	PAGKQTQQQRQQFANDGRGEI
			1	1		WRKLIKPNPRRQKS*CSTDPRQ
				ļ		KSTLISQREAVIRRFSEAGNFH
24712	55080	Α	24850	595	879	RRCSTTNIIKKIQQDKKGNRKA
		1	1	1		VHRSRFVHGRPITLLFFQLSGRI
1	1	I		1		QSHRLPGRAALFAGIPAACDQS
1		1	1			PSLRQPISSEWSDGR*HRRIFRS*
l				1		SEEF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24713	55081	Α	24851	1009	1478	IYSLSVSVWAGATVMESPVWM
						PIASKFSIEQTMMQLSFLSRTTSI
						SYSFOPISDSSISNSLVGERSRPR
						AQISSNSSRL*AIPPPLPPMVKD
						GRIMHGKPTSAATASASSIVCAI
						PERAVSR PIFFIATSKRRRSSALS
						IASAVAPIMVTPNSSNTP
24714	55082	Α	24852	48	209	GLFQVGYAENCRDNNHSFSLH
						SVVIGSSSERVVAKK*PDADHF
						SASFVPVILK
24715	55083	A	24853	1358	3194	PTASFAATLAIGKPVAFDASAE
					\	ERDTRGFISITIRRPFFGFTANVT
						RLETERGGICSRSVPWEVVSKI
1						AWAEAPDPPGVTILGLKTQHAT
1						/VSERVTSRFTRSHYRQFDLDQ
		1				AFSAKIFDRYLNLLDYSHKRKT
ł						ELGDEAAFSAKLDVFYDLYNL
İ		1				AQKRRFERYQYALSVLEKPMD
l						FTGNDTYNLDRSKAPWPKNEA
					ļ.	ELNALWDNVFSLAMTAFAREI
	1					DPHTNYLSPRNTEQFNTEMSLS
1						LEGIGAVLQMDDDYTVINSMV
		1				AGGPAAKSKAISVGDKIVGVG
						QTGKPMVDVIGWRLDDVVALI
						KGPKGSKVRLEILPAGKGTKTR
						TVTLTRERIRLEDRAVKMSVKT
1						VGKEKVGVLDIPGFYVGLTDD
		1				VKVQLQKLEKQNVSSVIIDLRS
						NGGGALTEAVSLSGLFIPAGPIV
		1				QVRDNNGKVREDSDTDGQVFY
		1		1		KGPLVVLVDRFSASASEIFARD
				1		VVVGLCICLSSLDVWLCGLFCR
		1		i		CSVDAERTKRQGKKGEESRKR
						RRKGYELSYLLFGWSAVVLLF
		1		l	1	VSDAEQHRASRHGGNACQQHP
				1		DVRHDEADGYPRQSHQLRLRR
			1	1		FRCFRPGRPLRLRRCQHERHDL
2.54	55001	<b>!</b>	0.1051		CON	PDDCRQVDRRRNGDWRGDDA
24716	55084	A	24854	100 852	671 1335	YFCFWTYPLLFPVCMDRKAGK
24/1/	22082	l <sup>A</sup>	24833	032	1333	HG\APAGDLYVOVOFNSHPIFE
						REGNTLYCEVPINFAMAALGGE
1			l	1		IEVPTLDGRVKLKVPGETQTGK LFRMRGKGVKSVRGGAQGDLL
1		1	1	1		
1				1		CRVVVETPVGLNERQKQLLQE
1		l		1		LQESFGGPTGEHNSPRSKSFFD
24710	55006	<del>  -</del>	24056	326	471	GVKKFFDDLTR PLOTHPAPQQRGHPIYQHGQTV
24718	55086	A	24856	320	4/1	
				1		S*SCRRRGWSSISRPVIHHADAQ
	1				L	LMP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nueleotide insertion)
1				sequence		
24719	55087	Α	24857	1095	1283	SCRRRGWSSISRPVIHHADAQL
						MP*PQIVYQHLKPPGIYRQQDR
						L*IPD*RPVLTIQDVRGP
24720	55088	Α	24858	1	1159	
24721	55089	Α	24859	1744	3019	KISPRRPQET/CFQKSTNIAREKT
		1				LEQEFSVQLFEKIGRRMCLTRE
		1				GKKLLPHIYELTRVMDTLREAA
1						KKESDPDGELRVVSGETLLSYR
1						MPQVLQRFRQRAPKVRLSLQSL
l						NCYVIRDALLNDEADVGVFYR
1						VGNDDALNRRELGEQSLVLVA
						SPOIADVDFTEPGRHNACSFIIN
		1				EPOCVFROIFESTLRORRITVEN
						TIELISIESIKRCVAANIGVSYLP
						RFAVAKELECGELIELPFGEOSO
1	1			ŀ		TITAMCAHHAGKAVSPAMHTFI
						HALYSCRMRRERLIRATRSCOF
l	1					NILHEPRRADKRSASGSLRLOE
	1					FNYODKPCMEIRIRTYEKTPIR
						MDKGLTNFGSRWILPELAHKE
	1	1				KDEETOORRKSKRDSRREETDT
						KKKTRDKATKREKDKEIEEMA
						SISITSRCEDKLENFSSMKYYTEI
24722	55090	IA	24860	321	575	SYRRYVFPAO*NRHLOFAVKPP
		1				APVIVHPTLVDSEHHHSLHDKR
	1					ROHRPHHSVAHHGSRNSSATA
						TINALLGRVAONAATRGTR
24723	55091	A	24861	935	1637	POPPPPLRREGAGLEKELGGRR
		1		1		VRVKGSGCOPWICGKVPVFIPV
	1					ITEOESFOELI/RRTTTACESLGK
	1				}	EYEILMIDDGSSDNSAHMLVEA
	1			l		SOAENSHIVSILLNRNYGOHSAI
	1					MAGFSHVTGDLIITLDADLONP
	1					PEEIPRLVAKADEGYDVVGTVR
				l	1	QNRQDSWFRKTASKMINRLIQR
			1			TTGKAMGDYGCM/LRAYRRHI
		1		l		VDAMLHCHERSTFIPILANIFAR
						RAIEIPVHHAEHS
24724	55092	A	24862	1020	1229	KAILII VIIIALII3
24/24	33092	ΙΛ.	24002	1020	1227	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequenc <b>e</b>		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24725	55093	Α	24863	98	1242	MDKFRVQGPTKLQGEVTISGA
						KNAALPNLFAALLAEEPVEIQN
						VPKLKDVDTSMKLLSQLGAKV
l						ERNGSVHIDARDVNVFCAPYD
				ì		LVKTMRASIWALGPLVARFGQ
						GQVSLPGGCTIGAHIVMDKVSV
						GATVTIMCAAILTEGATIIEVFY
		1.01				FLPLAIGKFNELYPNIKVRIL\EQ
1						GTNNCM\ESVLCN\ESDFG\INM\
						NNVTNSSI\DFTPL\VORRPFVL\
						ACRRDHPLAKKOLVEWOELVG
						YKMIGVRSSSGNRLLIEOOLAD
				i		KPWKLDWFYEVRHLSTSLGLV
1						EAGLGISALPGLAMPHAPYSSII
1						GIPLVEPVIRRTLGIIRRKDAVLS
						PAAERFFTLLSDFCPGVPSEDNI
1						FESPASOOHEYIGSROCFLMPSE
1	İ					ттотмну
24726	55094	A	24864	1	833	MAELGKRSSVAVSWIGKVKTT
				[		AQMVALAWLLWRPPIWVEYA
1			ŀ			GIALFFVAAVLTLWSMLQYLTL
						RVOICLISDRFGVIFSKRSKVVK
-						NIVDSSRQEYLEEFYARYNVELI
İ			ŀ			RAPEGFFYLRPRSTTLIPRSVLS
						ELDMMVGKILCYLYLSPERLA
						NEGIFTQQELYDELLTLADEAK
			ļ			LLKLVNNRSTGSDVDRQKLQE
						KVRSSLNRLRRLGMVWFMGH
						DSSKFRITESVFRFGADVRAGD
						DPREAQRRLIRDGD/AMPIENHL
						QLNDETEENOPDSGEEE
24727	55095	Α	24865	819	1428	SPRGISRGGCRCQPISGDG\RIFA
1			İ	1		GILHGLDNELPLQEEVEGNGLE
				1		QEGLPFPIRQSDALDLCFSDILV
			l	1		SPNMYIDDGMKRDWPVMKQC
			ŀ	1		ALQRRWRILPDHYGTEPAEKRP
				1		GFAPHYCFSYALLVHMMAQQC
			ĺ			DLEVGDFVWTGGDTHLYSNH
1			1			MDQTHLQLSREPRPLPKLIIKRK
			1			PESIFDYRFEDFEIEGYDPHPGIK
24728	55096	A	24866	2	1445	
24729	55097	Α	24867	I	1785	
				l		

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
}	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
		<u> </u>				
24730	55098	Α	24868	3	654	VRFFKGRGSSWHMV\SQWDYV
						ADRFGTADLLPFTISDMDFATA
1		l				PCIJEALNQRLMHGVFGYSRWK
1		ŀ				NDEFLAAIAHWFSTQHYTAIDS
1	1	ŀ	1			QTVVYGPSVIYMVSELIRQWSE
1	i					TGEGVVIHTPAYDAFYKAIEGN
		ŀ				QRTVMPVALEKQADGWFCDM
1		1			1	GKLEAVLAKPECKIMLLCSPQN
1						PTGKVWTCDELEIMADLCERH
		l				GVRVISDEIHMDMVWGEOPHIS
1		1			1	LE
24731	55099	A	24869	955	1278	TLTPDFL*QGHFFRFAGIAYQIV
						KOTTVGTTVEITLRRYVVDGNR
		ı				IPPRODROGDAFFGFATHTOOW
Į.						HQVFKRQRDIQIVTANTATTVT
		l				QQAGFTIETRTPLRPHQQQ
24732	55100	A	24870	<del> </del>	1707	QQAGITIETKITEKITQQQ
24732	55100	Ā	24871	i	3615	
24734	55101	A	24872	i -	1931	MLLLREHTDTRHDHPRTPTRAT
24/34	33102	Γ.	24072	1'	1991	RRRRHTEKKRRKKKRPATHVL
						MTATLIPVSRLLATDQPQLFHE
ı	1	l l				SASKPAPHLVSSLGCHCGDASC
1	1	ļ				SGRAVADAMOFKTVFPVFVAA
	ļ	ı				
1		l				AMNPEHMAESGHRITRPEFPDY
		1			1	RELFRESDIKSAVVDSICQCNTP
1		l				FNYLFRCFELQCLSRHGHSAYG
		ł				VMQRRHSPMQETRIASLPLQRY
1		l				RWHEDSVRNRWFSVMVGPSVR
						VNEWFSAYAMAGMAYSRVST
						FSGDYLRQPLVSGFHEA
24735	55103	Α	24873	1281	2234	ISTNSILRIFSGFRKLMEV/TRLR
		1				DTPILTFMNKLDR/DIRDPMELL
		1				DEVENELK/IGCAPITWPIGCGK
		l			1	LFK/GVYHLYKDETYLIQEVRIV
					į	KGLNNP/DLDAAVGEDLAQQL
	ļ	1				RDE/LELVKGASNEFDKELF/LA
	1	l				GEITPVFFGTALGNF/GVDHML
						DG\FMAGDRSHVEEAY/PGDIL
1	1	1	1		I	G\QGEMMKFTGIPNF/APELFRR
		1				IRLKDPLKQK/QLLKGLVQLSEE
l		l			1	GA VQV/FRPISNNDLIVGA VGV
I	1	1			1	LOFDVVVARLKSEYNVEAVYE
1	1	1	1		1	SVNVATARWVECADAKKFEEF
		1			1	KRKNESQLALDGGDNLAYIATS
1					1	MVNLRLAQERYPGRQFHQTRE
		ഥ_		L		I TILLLA QUALITURQI AQIAE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24736	55104	A	24874	2	760	AKFEEKFLAVPAEALVYTMKG
						DQKYFPVYANDGKLLPNFIFVA
						NIESKDPQQI/DRIQALAGWIAE
						Q\IGADVNHATRAGLLSKCDLM
						TNMVFEFTDTQGVMGMHYAR
						HDGEAEDVAVALNEQYQPRFA
						GDDLPSNPVACALAIADKMDT
				l		LAGIFGIGQHPKGDKDPFALRR
						AALGVLRIIVEKNLNLDLQTLT
						EEAVRLYGDKLTNANVVDDVI
		1				DFMLGRFRAWYQDEGYTVDTI
l				-		QAVLARRPTRPADFDDWHA
24737	55105	A	24875	2	391	
24738	55106	A	24876	2	1785	WLLYRAHVKGE/STEALLPNM
						VATSLAKLPIPKLMRWGATYV
					İ	HFVRPVHTVTLLLGDKVIPATIL
		l			ŀ	GIQSDRVIRGHRFMGEPEFTIDN
			ŀ			ADQYPEILRDSGKVIADYEERK
						AKIKADAEEAARKIGGNADLSE
		l	ŀ			TLLEEVASLVEWPVVL/TAKF/E
		1				EKF/LAISRTALVPYSADNMYQ
1	l	l				LVNDVQSYPQFLPGCTGSRILES
İ						TPGQMTAAVDVSKAGISKTFTT
į						RNQLTSNQSILMNLVDGPFKKL
1		1	l			IGGWKFTPLSQEACRIEFHLDFE
		1	Ì		}	FTNKLIELAFGRVFKELAANMV
İ		1	Ì		!	QAFTVRAKEVYSASMRYIYVIV
			İ		1	YISYRPPNLLDSHIIGFSSITTHC
						AVTSHHYNSCDMTMTIHSYVT
						PYISVRVSLTQTYAPSPSRNLRI
						TDLTIYRAISSEKSIDDTRHIHTI
						ESRCRVCGAHDSRDAERLDRF
		l	l			AQTADEIRIADRGFSSRPECIRS
		1				LAFGEADYIVRVHWRGLRWLT
1			1	I	l	AEGMRFDMMGFLRGLDCALIS
1		l	l	I	l	KTRLLSENRRKGRVVQAETLE
						AAGHVLLLTSLPEDEYSAEQVA
1		1	l			DCYRLRWQIELAFKRLKSLLHL
		1				DALRAKEPELAKVGIFANLLAA
		١		ļ		FLIDDIIQPSLDFPPRSAGSEKKN
24739	55107	A_	24877	1	2022	
24740	55108	A	24878	2344	2523	
24741	55109	Α	24879	82	732	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24742	55110	A	24880	828	2184	SGCILPPESLRLWASVARRPRTP NOPROGERIGISPAGSALALA CGKRTVLFASLCTRTGGSQLGE PALVLLADQMLALDDQYTFLQ RIPRSPYGLAFOPVPHLHYLING MRLLDLAPIAFRAMSLAFPVNY LDAVRTGSICSNCNQRSNDSAM GISMNYSTLMRVRAIWVYAAY MERVLDNLLNNALRYCHSTVE TSLLLSGNRATLIVEDDGPGIAP ENREHIFEPFVRLDPSRDRSTGG CGLGLAIVHSIARGNWATSDHR SLTGTQPYARIVVFLVLIFAFR CYDLTGLYSFVFRVVVQNAFPG CPLGIGDVHYYAVAYGVFD LSSRTGTTVFFDDLNTPDFDGC CFLGGGVHYYAVAYGVFD RAMAYGERERFQLYRRGRY VEFNLPSELAATVYQSDPNFAT GCFRLYTSCNTFNMHKAATML
24743	55111	A	24881	679	1628	VRAEFSQGYOHQV PRADARKAYDTICLAFGCDAOF IEITFTRLPEVRIHHLQSPASALS SCNHLLIRITOFNADTIGHHLP VNRRVYTVLRDMRHHGDIVF GVRRGVEHYAAMDPGIVKEIM EVRLLFLAVFVGNHHSRNDSLP VQFVVHTNGDTYTFPGNNIRRD IGFKWRIAALMLHHFLIIDPHFA VMSDRIETQHDTLACPHFRYKD KAPVYRWYHGYKALSKEIL VYQRLYQIFOHAEYQTGTNDA EISTKLCLFSSSLLLLSSSSLOHSI PPNACL
24744	55112	A	24882	1	1515	TIMES
24745	55113	Α	24883	1954	2859	
24746	55114	A	24884	121	393	SLARWIICNTSEELAANIKRLGD VTVRLNAFHAHGVYGEAQSM GERHAGRR*AVG*CVKGARGE PVDISGAQL/RGCGVNALSGLR FGTDL

SEQ ID	SEQ ID NO:					Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24747	55115	Α	24885	1	3706	MDLGANGWQTFRYVVLPNLSS
		1				ALLAGGMLAFALSFDEIIVTTFT
	ŀ	1		l		AEQVDAAVRAADAAFAEWGQ
				ŀ	l	TTPKVRAECLLKLADVIEENGO
						VFAELESSNNTAFARASSNGDL
					1	PTKADLQAQLDSLNKQKDLSA
				1		ODKLVOODLTDTLATLDKIDRI
1		İ				KEETVQLRQKVAEAPEKMRQA
	l					TAALTALSDVDNDEETRKILST
İ			1			LSLRQLETRVAQALDDLQNAQ
				1		NDLASYNSQLVSLQTQPERVQ
						NAMYNASQQLQQIRSRLDG
24748	55116	Α	24886	2842	5034	NVRLSPAMRCRWRRGSPMPAM
				F		FEVGVKTRIIORSREVILMADHS
	l			1		KFDAVEPHAVATLSCIKTIISDS
		ł		1		GLPETIAORYORAGSSLPEAOP
		l		1		DREIEKRGPAIAQAFDAEGKPS
		ļ.	1	ŀ		KASEGWARGCGITVDQAERLT
1		1				TDKGEWLLYRAHVKGESTEAL
		ŀ				LPNMVATSLAKLPIPKLMRWG
				İ		ASDVHFV\WPVHTVTLL\LGDK
		1		1		VIPATILGIOSDRVIRGHRFMGE
	ł	1				PEFTIDNADQYPEILRERGKVIA
						DYEERKAKIKADAEEAARKIGG
						NADLSESLLEEVASLVEWPVVL
		1		1		TAKFEEKFLAVPAEALVYTMK
1		ı		l .		GDOKYFPVYANDGKLLPNFIFV
		ı		l		ANIESKOPOOIISGNEKVVRPRL
		ı		1		ADAEFFFNTDRKKRLEDNLPRL
		1				OTVLFQQQLGTLRDKTDRIQAL
				i		AGWIAEQSGTDAGDQTGYSDE
				ı		RWVPDAHSEAPSPVSDMLSAL
1						LLLTLVLAGMVRMAARLLMA
		1				KPPOAVNRGDIGWLTTSPMVIL
		1	1			LVMMLAMGTHIPOPVIRIPGGR
1		1				FHYSPHOGRTDLPAQRSTWHD
		1		i		FCLOAPHLFRRNTVNVNSSSNR
1		1				GEAILAALKTOFPGAVLDEERO
	1	1		1		TPEQVTITVKINLLPDVVQYLY
1						YOHDGWLPVLFGNDERTLNGH
1	i	1		1		YAVYYALSMEGAEKCWIVVK
ł	1			I		ALVDADSREFPSVTPRVPAAV
1	1	1		1		WGEREIRDMYGLIPVGLPDORR
				1		LVLPDDWPEDMHPLRKDAMD
24749	55117	A	24887	49	531	
24750	55118	Α	24888	1619	1968	
24751	55119	Α	24889	1	553	
24752	55120	Α	24890	744	970	LAVISEASNQKRGDPVDHAD*I
1		1				AVANMLLNQFQQSYTPMLHDL
	1	1	1			IVRYDGDIHQQKSFGVQLPGKL
		L				RIVKLIQPVE

	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
24753   55121   A   24891   52   455	NO:		hod				
24753   55121		sequence	1	09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
24754   55122		1			sequence		
24754   55122	24753	55121	A	24891	52	455	
FHSEDVYRHPVARIVNAYEA WEEAPQKTMTPANQHAGGLV QCDKMLQRSREVIELSEQN, AQMRHVRGADMAMIFQEPMT SLNPPFTVGEQLAESIRLHQNA: REEAMVEAKRALDQVRIPEAC TILISRYPPQLSGGMRQRVMIA MALSCRPAVLIADEPTTALDVY IQAQILQLIKVLQKEMSMGVIF THDMGVVAEIDARVLVMYQG EAVETGTVEQIFHAPQHPYTRA LLAAVPQLGAMKGLDYPRFFF LISLEHPAKQAPPIEQKTVVDG PVLRVRNLVTRPPLRSGLLNR: TREVHAVEKVSPDLWGETLS VGESGSKSTTGRALLRLVESC GGEIIFNQQRIDTLSPGKLQALE RDIQFFQDPV3ASLDPRGTIGDS IEPLRVHGLLPGKDAAARVAW LLERYGLLPEHAWRYPHEFSG QRQRICIARALALNFVIIADE/ VVSALDVSIRGQIINLLLDLQRDI GIAYLFISHDMAVVERISHRVA VMYLGQIVEIGPRRAVENPOP PYTRKILLAEPPVAEPSQRPQ VLLSDLPSNIHLRGEAVVAL KPGAQIGRIPQSWT IRTGRNESYKGDISRYKFVSAR ALRNEVRGGRCHKTIRRVSVG KKVIGLKFRILVPVVTPESIVP PAPVANDTLVAEVSDAPQANE PTFNNEDLA*FAVLSTPTAILIS PGRWPAAGSECVGRGHAH ELALRAFSCCCFSSNAISIPN TDAPGDEEASLQRAQAQGVGH IIVPATEAENFARVLALAENNYQ PLYAALGLHRGMLEKHSDVSL GQAERFVQLGYKIGVGGTTTYP ASKTRDVAKLPLASLLETLD PMPLNGFQGGPORPEQAARN HLKRHDPRRGVVYGFGSGLQ QAERFVQLGYKIGVGGTTTYP ASKTRDVAKLPLASLLETLD PMPLNGFQGGPORPEQAARN FAVLCELRREPADEIAQAVLN	24754				2988	3614	PVSVLTQKRLSPATSHRSTCRV
WERAEQKTMTPANGHAGGLV QCDKMILQRRSREVIELSEQN. AQMRHVRGADMAMIFGEPMT SLNPVFTVGEQLAESIRLHQNA: REEAMVEAKRINLDQVRIPEAC TILSRYPHOLSGGMRQRVMIA MALSCRPAVLIADPTTALDVT IQAQILQLIKVLQKEMSMOVIF THDMGVVAEIADRVLVMYQG EAVETGTVEGIFHAPQHPYTRA: LLAAVPQLGAMKGLDYPRFFE LISLEIPAKQAPPIEQKTVVDG PVLRVRNLVTRPPLRSGLLNRX TREVHAVEKVSPDLWPGFTLS VGESGSKSTTGRALLELVESS GGEIFINGQRIDTLSPGKLQALF RDIOFIFODPYASLDPROTIGOS IEPLRVHGLLPGHAWRYPHEPSG QRQRICIARALAINPKVIIADE VSALDVSIRGQINNLLDLQRDI GIAVLFISHDMAVVERISHRVA VMYLGQIVEIGPRRAVFENPQI PYTRKILLAEDPVAEPSRQRPQ VLLSDDLPSNIHRIRGEAVVAL KPGAQIGRPQSWT RAVINGRESS ALRIEVRGGRCHKTIRRVSVG KKVIGLKFRRILVPVVTPESIVIF PAPVANDTLVAEVSDAPQANDE PTFRNEDLA*FAVLSTPTAILIS PGRWPAAGSECVIKGAHAH ELALRAFSCCCFSSNAISIPN TDAPGDEEASLQRAQAGVGR IVPATEAENFARVVAVGEIG DLFGDDPQFERQOWLLDEQLII VVKRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERRYQLGYKIGVGGTTTYP ASKTRDVAKLPLSLLELTED PDMPLNGFQGQPNREQAARN FAVLCELRREPADEIAQAVLNN TYTLVNNP		ŀ	l			ļ.	LNOAYVTPSKCWPDVEEISFNF
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GGEIIFNGQRIDTLSPGKLQALF RDIQFIFQDPYASLDPRQTIGDS IEPLRVHGLLPGKDAAARVAW LLERVGLLPEHAWRYPHEFSG QRQRICIARBALAINPEVIIADE VSALDVSIRGQIINLLLDLQRDI GIAYLFISHDMAVVERISHRVA VMYLQQIVEIGPRAAVFENPGI PYTRKILLAEDPVAEPSRQRPQ VLLSDDLPSNIHLRGEGAVVAL KFGAQIGRPQSWT IRTGRNESVKGDISR VKFVSAR ALRNEVRGGRCHKTIRRVSVG KKVIGLKFRHLVPVVTPESIVIF PAPVANDITLVAEVSDAPQANDE PTFRNEDLA*FAVLSIPTAILIS PGRWPAAGSCOVGGAHAH ELALRAFSCCCFSSNAISIPN TDAPGDEEASLQRAQAGVGR IIVPATEAENFARVLALAENTV PLYAALGLHFGMLEKHSDVSL GOLQQALERRPAKVVAVGGIG DLFGDDPQFERQQWLLDEQLII VVKRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERTVQLGYKIGVGGTITYPR ASKTRDVAKLPLASLLLETIA PDMPLNGFQGQPNRPQAARVA FAVLCELRREPADEIAQAVLNI TYTLVNVP	ŀ						TREVHAVEKVSFDLWPGETLSI
ROJOFIFODPY ASL DPROTIGOS IEPLRVHGLLPGK DAAARVAW LLERVGLLPEHAWRYPHEPSG QRQRICIARALALNPKVIIADE VSALDVSIRGQINILLDLQRDI GIAYLFISHDMAVVERISHRVA VMYLQQIVEIGPRAAVENPOP PYTRKILLAEDPVAEPSQRPQ VLLSDDLPSNIHLRGEEAVVAL KPGAQIGRPPOSWT IRTGRNESYKGDISRYKFVSAR ALRNEVRGGKCHHTIRRVSVG KKVIGLKFRILLPVVTTESIVIP PAPVANDTLVAEVSDAPQAND PTFNNEDLA*FAVLSTPTAILIS PGRWGPAAGSECGVRGAHAH ELALRAFSCCCFSSNAISIPN 1444 TDAPGDEEASLQRAQAGVGVG IIVPATEAENFARVLALAENYQ PLYAALGLHGMLEKHSDVSL EQLQQALERRPAKVVAVCGIG DLFGDPPOFERQQWLLDEQLII VKRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERVQLGYKIRGVGGTTTYPR ASKTRDVIKALPLASLLETDA PDMPLNGFQGPORPREQA AR FAVLCELRREPADEIAQAVLNN TYTLVNNP			1				VGESGSGKSTTGRALLRLVESQ
IEPLR VHÖLLPGKDAAARVAW LLERVÖLLPEHA WRYPHEFSG QRQRICIARALALNPKVIIADE- VSALDVSIRGQIINLLLDLQRDI GIAVLFISHDMAVVERISHRVA VMYLGGIVEIGPRRAVEENPOP PYTRKILLAEDPVAEPSRQRPQ VLLSDDLPSNIHLRGEEAVVAL KPGAQIGRPQSWT IRTGRNESVKGDISR VKEVVSAR ALRNEVRGGRCHKTIRRVSVG KK VIGLKFRHLVPVVTPESIVIF PAPVANDTLVAEVSDAPQANE PTFNNEDLA*FAVLSITTAILIS PGRWGPAAGSECVIRGAHAH ELALRAFSCCCSSNAISIEN TDAPGDEEASLQRAAQAGVGR IIVPATEAENFARVLALAENVQ PLYAALGLHPGMLEKHSDVSL EQLQQALERRPARVVAVGEGE DLFGDPPOFERQOWLLDEQLII VVKRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERTVQLGYKIGVGGTITYPR ASKTRDVAKLPLASLLEETDA PDMPLNGFQGGPNRFEQAARV FAVLCELREPADEIAQAVLN TYTLVNNP							GGEIIFNGQRIDTLSPGKLQALR
LLERVGLIPEHAWRYPHEFSG QRQRICIARALALNPKVIIADE QRQRICIARALALNPKVIIADE VSALDVSIRGQINLLLDLQRDI GIAVLISHDMAVVERISHRVA VMYLGOVEIGPRAVPENPO PYTRKILLAEDPVAEPSRQRPQ VLLSDDLPSNIHLRGEEAVVAL KPGAQIGRPQSWT LKTGRNESVKGDISRYKFVSAR ALRNEVRGGRCHKTIRRVSVG KKVIGLKFRHLVPVVTESIVIE PAPVANDTLVAEVSDAPQAND PIFFANEDLA*PAVLSTPTALLIS PGRWGPAAGSECGVRGAHAH ELALRAFSCCCFSSNAISIFN 1444 TDAPGDEEASLQRAAQAGVG IIVPATEAENFARVLALAENVQ PLYAALGLHPGMLEKHSDVSL EGOLQQALERRPAKVVAVGEIG DLFGDDPOFERQOWLLDEQLII VKKYELPVILLARSRTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERVQLGYKIGVGGTTTYP ASKTRDVIKALPLASLLETDA PDMPLNGFQGQPNRFEQAAR FAVLCELRREPADEIAQAVLN TYTLVNVP			1				RDIQFIFQDPYASLDPRQTIGDS
QRQRICIARALALNPKVIJADE/ VSALDVSIRGQIINLLIDLQRDI GIAYLFISHDMAVVERISHRVA VMYLGGIVEIGPRRAVERISHRVA VMYLGGIVEIGPRRAVERINOP PYTRKILLAEDPVAEPSQRPQ VLLSDDLPSNIHLRGEEAVVAL KRGAQIGRYQSWT IRTGRNESYKGDISRYKFVSAR ALRNEVRGGKCHKTIRRVSVG KKVGLKFRILVPVVTPESIVE PAPVANDTLVAEVSDAPQANE PTFNNEDLA*FAVLSIPTAILIS PGRWGPAAGSECVRGAHAFI ELALRAFSCCCFSSNAISIPN TDAPGDEEASLQARAQAGVG PLYAALGLHRGMLEKHSDVSL EQLQQALERRPAKVVAVGEIG DLFGDPPQFERQQWLLDEQLI VKRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYGFGSGL QAERFVQLGYKIGVGGTTTYPR ASKTRDVIAKLPLASLLETIDA PDMPLNGFQGGPNRPEQAARV FAVLCELRREPADEIAQAVLN	1		1				IEPLRVHGLLPGKDAAARVAW
VSALDVSIRGQINLLLDLQRDI GIAYLFISHDMAVVERISHRVA VMYLQQIVEIGPRAVFENPGI PYTRKILLAEDPVAEPSRQRPQ VLLSDDLPSNIHLRGEAVVAU KFGAQIGRPQSWT  24755 55123 A 24893 134 595 IRTGRNESYKGDISRYKFVSAR ALRNEVRGGRCHKTIRRVSVG KKVIGLKFRHLVPVVTPESIVIE PAPVANDITLVAEVSDAPQANDE PTFRNEDLA*PAVLSIPTAILIS PGRWPAAGSECVIKGAHAH ELALRAFSCCCFSSNAISIPN  708 1444 TDAPGDEEASLQRAQAGVGR IIVPATEAENFARVLALAENVY PLYAALGLHRGMLEKHSDVSL EQLQALERRPAKVVAVGEIG DLFGDDPQFERQQWLLDEQLII VVKRYELPVILHSRKTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERTVQLGYKIGVGGTITYPR ASKTRDVAKLPLASLLLETIDA PDMPLNGFQGQPNRPEQAARV FAVLCELRREPADEIAQAVLNI TYTLVNVP	1						LLERVGLLPEHAWRYPHEFSGC
GIAVLFISHDMAVVERISHRVA	1	1	1				QRQRICIARALALNPKVIIADEA
VMYLGOJVEIGPRRAVERNOH PYTRKILAEDPVAEPSRQPQ VULSDDLPSNIHLRGEEAVVAL KPGAQIGRPQSWT RTGRNESYKGDISRYKEFVSAR ALRNEVRGGRCHKTIRRVSVG KKVIGLKFRHLVPVVTPESIVIF PAPVANDTLVAEVSDAPQANE PTFNNEDLA*FAVLSITTAILIS PGRWGPAAGSECVIRGAHAH ELALRAFSCCCSSNAISIPN TDAPGDEEASLQRAAQAGVGR IIVPATEAENFARVLACHSDVSL EQLQALERRPARVVAVGEIG DLFGDPPOFERQWLLDEQLII VVKRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERTVQLGYKIGVGGTITYPR ASKTRDVAKLPLASLLEETDA PDMPLNGFQGGPNREQAARV FAVLCELRREPADEIAQAVLNN TYTLVNVP					1		VSALDVSIRGQIINLLLDLQRDF
PYTRKILLAEDPVAEPSRQRPQ VLLSDDLPSNIHLRGEAVVAL KFGAQIGR:PQSWT 24755 55123 A 24893 134 595 IRTGRESVKGDISR:VKFVSAR ALRNEVRGGRCH:TIRRVSVG KKVIGLKFRHLVPVVTPESIVIF PAPVANDTUVAEVSDAPQAND PTFNNEDLA*PAVLSTPTAILISI PGRWGPAAGSECGVRGAHAH ELALRAFSCCCFSSNAISIPN TDAPGDEEASLQRAAQAGVGW IIVPATEAENPARVLALAENVQ PLYAALGI-HPGMLEKHSDVSL EQLQQALERRPAKVVAVGEIG DLFGDDPQFERQWLLDEQLII VKKYELPVILLSKRTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERVQLGYKIRGVGTTTYPF ASKTRDVIAKLPLASLLETDA PDMPLNGFQGQPNRPEQAARV FAVLCELRREPADEIAQAVLNN TYTLVNVP							GIAYLFISHDMAVVERISHRVA
VLLSDDLPSNIHLRGEEAVVAL KPGAQIGRPQSWT 24755 55123 A 24893 134 595 IRTGRNESYKGDISRYKFVSAR ALRNEVRGGRCHKTIRRVSVG KKVIGLKFRILLPVVVTPESIVIP PAPVANDTLVAEVSDAPQANE PTFNIEDLA*FAVLSTPTAILLS PGRWGPAAGSECCVRGAHAH ELALRAFSCCCFSSNAISIPN 10444 TDAPDEEASLGAAQAGVG PLYAALGLHRGMLEKHSDVSL EQLQQALERRPAKVVAVCEIG DLFGDPPQFERQQWLLDEQLII VKRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYGFGSLQ QAERVQLGYKIRGVGGTTTYPR ASKTRDVIAKLPLASLLEETDA PDMPLNGFQGGPNRPEQAARV FAVLCELRREPADEIAQAVLN TYTLVNVP	1					1	VMYLGQIVEIGPRRAVFENPQH
24755 55123 A 24893 134 595 IRTGRNESYKGDISRYKFVSAR ALRNEVRGGRCHKTIRRVSVG KKVIGLKFRHLVPVVTPESIVIE PAPVANDTLVASVSDAPOAND PTFNNEDLA*FAVLSIPTAILIS PGRWGPAAGSECVGKGHAH ELALRAFSCCCFSSNAISIPN 11444 TDAPGDEEASLQRAQAGVGR IIVPATEAENFARVLALAENVQ PLYAALGLHRGMLEKHSDVSL EQLQALERRPAKVVAVGEIG DLFGDDPQFERQQWLLDEQLII VVKRYELPVILHSRKTHDKLAM HLKRHDLPRTGVVGFSGSLQ QAERTVQLGYKIGVGGTITYPR ASKTRDVAKLPLASLLLETDA PDMPLNGFQGPNRFQAARVA FAVLCELRREPADEIAQAVLNN TYTLVNVP							PYTRK\LLAEDPVAEPSRQRPQF
24755 55123 A 24893 134 595 IRTORNESVKGDISRYKEVSAR ALRNEVRGGRCHKTIRRVSVG KKVIGLKFRHLVPVVTESIVIE PAPVANDTLVAEVSDAPQAND PTFNNEDLA*PAVLSTPTALLISL PGRWGPAAGSECGVRGAHAHI ELALRAFSCCCFSSNAISIPN TOACH PAPVANDTLVAEVSDAPQAND PLYAALGALPROMALALAENVQ PLYAALGALPROMALAENVQ PLYAALGALPROMALALAENVQ PLYAALGALPROMALALAENVQ PLYAALGALPROMALAENVAVCEGE DLFGDDPOFERQOWLLDEQLII VKKYELPVILARSTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERVQLGYKIGVGGTTTYPR ASKTRDVIKALPLASILLETDA PDMPLNGFQGQPNRPEQAARV FAVLCELRREPADEIAQAVLNYTYTLVNVP	i		1				VLLSDDLPSNIHLRGEEAVVAL
ALRNEVRGGRCHKTIRRVSVG KKVIGLKFRHLVPVVTPESIVIF PAPVANDTLVAEVSDAPQANE PTFNNEDLA*FAVLSIPTAILIS PORWGPAAGSECOVRGAHAH ELALRAFSCCCFSSNAISIPN TDAPGDEEASLQRAAQAGVGR IIVPATEAENFARVLALAENVQ PLYAALGLHFGMLEKHSDVSL EQLQQALERRPAKVVAVGEIG DLFGDPOFERQOWLL DEQLII VVKRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERVQLGYKIGVGGTITYPR ASKTRDVAKLPLASLLEETDA PDMPLNGFQGPNRPEQAARV FAVLCELRREPADEIAQAVLNN TYTLVNVP							KPGAQIGR\PQSWT
KKVIGLKFRHLVPVVTPESIVIE PAPVANDTUVAEVSDAPQANE PTFNNEDLA*FAVLSTPTALLISE PORWOPAAGSECGVRGAHAH ELALRAFSCCCFSSNAISIPN TDAPGDEEASLQRAAQAGVGR IIVPATEAENFARVLALAENYQ PLYAALGLHPGMLEKHSDVSL EQLQQALERRPAKVVAVGEIG DLFGDDPGFERQQWLLDEQLII VKKYPLEPVILBSRTHDKLAM HLKRIDLPRTGVVYGFSGSLQ QAERFVQLGYKIGVGGTTTYPR ASKTRDVIAKLPLASLLLETDA PDMPLNGFQGQPNRPEQAARV FAVLCELRREPADEIAQAVLN TYTLVNVP	24755	55123	Α	24893	134	595	IRTGRNESYKGDISRYKFVSAR
PAPVANDTLVAEVSDAPQANE PTFNNEDLA*FAVLSTPTAILIS PGRWGPAAGSECGVRGAHAH ELALRAFSCCCFSSNAISIPN 24756 55124 A 24894 708 1444 TDAPGDEEASLQRAAQAGVG PL-YAALGLHRGMLEKHSDVSL EQLQQALERRPAKVVAVCGEG DLFGDPPQFERQQWLLDEQLII VKRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYQFGSGLQ QAERVQLGYKIGVGGTITYPR ASKTRDVIAKLPLASLLEETDA PDMPLNGFQGQPNRPEQAARV FAVLCELRREPADEIAQAVLNN TYTLVNVP			1	1			ALRNEVRGGRCHKTIRRVSVG
PIFRNEDLA*FAVLSIPITÄLIS PGRWGPAAGSECGVRGAHAH ELALRAFSCCCFSSNAISIPN TDAPGDEEASLQRAQAGVG IIVPATEAENFARVLALGENYO PL.YAALGLHFGMLEKHSDVSL EQLQALERPRAVVAVGEIG DLFGDDPGFERQQWLLDEQLII VVKRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVGFGSGLQ QAERVQLGYKIGVGGTITYPR ASKTRDVAKLPLASILLETDA PDMPLNGFQGPNRFEQAARV FAVLCELRREPADEIAQAVLNI TYTLVNVP	1		l	1		1	
PGRWGPAAGSECGVRGAHAHI ELALRAFSCCGSSNAISIPN 24756 55124 A 24894 708 1444 TDAPGDEEASLQRAAQAGVG PLYAALGLHPGMLEKHSDVSL EQLQQALERRPAKVVAVGEIG DLFGDDPQFERQQWLLDEQLII VKKYFLEPVILHSRFTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERVQLGYKIGVGGTITYPR ASKTRDVIAKLPLASLLETDA PDMPLNGFQGQPNRPEQAARV FAVLCELRREPADEIAQAVLNY							PAPVANDTLVAEVSDAPQAND
24756 55124 A 24894 708 1444 TDAFGDEEASLQRAAQAGVGH IIVPATEAENIFARVINALAENIVQ PLYAALGLHFGMLEKHSDVSL EQLQQALERRPARVVAVGEIG DLFGDDPOFERQOWLLD EQLII VVKRYELPVILHSRRTHDKLAM HILKRHDLPRTGVVYGFSGSLQ QAERRVQLGYKIGVGGTITYPR ASKTRDVAKLPLASLLEETDA PDMPLNGFQGQPNRPEQAARV FAVLCELREPADEIAQAVLNN TYTLENNP							
24756 55124 A 24894 708 1444 TDAPGDEEASLQRAAQAGVGHIVPATEAENFARVLALAENYQPLYAALGLHRGMLEKHSDVSL EQLQQALERPAKVVAVOEIG DLFGDDPGERQQWLLDEQLII VVKRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERFVQLGYKIGVGGTITYPR ASKTRDVIAKLPLASLLLETDA PDMPLNGFQGQPNRPEQAARV FAVLCELRREPADEIAQAVLNY							PGRWGPAAGSECGVRGAHAHF
IIIVPATEAENFAR VLALA ENYQ PL YAALGHPGMLEKHSDVSL EQLQQALERRPAKVVAVCEIG DLFGDDPOFERQOWLLDEQLII VKK YEL PVILHSR THDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERRVQLGYKIGVGGTTTYPR ASKTRDVIAKLPLASLLETDA PDMPLNGFQGQPNRPEQAARV FAVLCELRREPADEIAQAVLNN TYTLVNVP							
PLYAALGLHPOMLEKHSDVSL EQLQQALERPAKVVAVGEIG DLFGDPPQFERQQWLLDEQLI V\KRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERFVQLGVKIGVGTTTYPP ASKTRDVIAKLPLASLLLETDA PDMPLNGFQGPPNRPEQAARV FAVLCELRREPADEIAQAVLNN TYTLVNVP	24756	55124	Α	24894	708	1444	
EQLQQALERPAKVVAVGEIG DLFGDDPGERQQWLLDEQLII VKKPYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYGFSGSLQ QAERFVQLGYKIGVGGTITYPR ASKTRDVIAKLPLASILLETDA PDMPLNGFQGQPNRPEQAARV FAVLCELRREPADEIAQAVLNY							
DLFGDDPQFERQQWLLDEQLII V\kryelpvilhsrrthdklam Hilkrhddprtgovygfsolg Qaerfvolgykigvggtitypb Asktrddvaklplasllletid pdmplngfqggpnrbeqaarv Favicelrepadeiaqavlnn tytlvnyp					l .		
V\KRYELPVILHSRRTHDKLAM HLKRHDLPRTGVVYGFGSSLQ QAERFVQLGVKIGVGGTTTYPB ASKTRDVIAKLPLASLLLETDA PDMPLNGFQGPNRPEQAARV FAVLCELRREPADEIAQAVLN TYTLVNVP			1				
HLKRHDLPRTGVVYGFSGSLQ QAERFVOLGVKIGVGGTITYPR ASKTRDVIAKLPLASLLLETDA PDMPLNGFQGQPNRPEQAARV FAVLCELRREPADEIAQAVLN TYTLVNVP	1		1				
QAERFVQLGYKIGVGGTITYPR ASKTRDVIAKLPLASLLLETDA PDMPLNGFQQPNRPEQAARV FAVLCELRREPADEIAQAVLNY TYTLVNVP			1		1		V\KRYELPVILHSRRTHDKLAM
ASKTRDVIAKLPLASLLLETDA PDMPLNGFQGPNRPEQAARV FAVLCELRREPADEIAQAVLN TYTLVNVP			1		l		HLKRHDLPRTGVVYGFSGSLQ
PDMPLNGFQGQPNRPEQAARV FAVLCELRREPADEIAQAVLNN TYTLVNVP			1				QAERFVQLGYKIGVGGTITYPR
FAVLCELRREPADEIAQAVLNN TYTLVNVP	1		1				ASKTRDVIAKLPLASLLLETDA
TYTLVNVP				1			PDMPLNGFQGQPNRPEQAARV
	1		1				FAVLCELRREPADEIAQAVLNN
24757 55125 A 24895 1460 1777							TYTLVNVP
	24757	55125	Α	24895	1460	1777	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24758	55126	A	24896	381	482	IPGPQSPQSSPDLGWLQC*NP*R LLWKAPRSDK
24759	55127	Α	24897	977	1144	
24760	55128	A	24898	2	280	CTFALRDEYPRMLNVDNKGLR DHIELQPLAV/DASYPGGDFRK NA VYVPMW*SAHCEPEGRFAE TGSRRYGEDFASTAPGFRRAAT RKRVRRKR
24761	55129	В	24899	1	1665	
24762	55130	Α	24900	296	495	
24763	55131	Α	24901	1	5352	
24764	55132	A	24902	187	280	AVDTGLADIIRD\EPNMVARLP LSFTASLSIS
24765	55133	A	24903	1775	1981	HIWQNSLIVLFRGCRSAHAKVH RWKN*LPLNLAPLLPRSGSSAPI RPPPSAQARQPMKSTYGVDRR HS
24766	55134	Α	24904	760	1473	
24767	55135	A	24905		2124	IMNRMSGYYWQLFAESGETCR ARELEQCLEDFYDQRKESVRW HLDLHTAIRGSLHPQFGVLPQR DIPWDEKFLTWLGAAGLEALV FHQEFGGTFHESARHFGALV FHQEFGGTFHESARHFGALV FMGERGFHFHSARHFGALV FMGERGFHFHSARHFGALV FMFEKGTLLAQDGEERTTVTH DVEVAKNATHEIKTOFEKEKLA QDRIKRNQ/LDANIQRLNYSLDI ANAAGIKEPYYSNDGIERKLEI EKAYTDVAELMGELRNIQVL EQLITKAHVNDVNFTPFKYQLSP SLPVKKDGPGKALIVILSALIGG MLVGRVCIKRNADSPEVFSGFS DPNVVLIAAVFIIGDGLVRTGV VLIAMVFILOGLVRTGV VLIAMVFILOGLVRTGV VNSELLBEGVHGFSFFSVTPIGL UVLVLGILIVML/VMFFMLKGDT QTPQREGWTRTFRDLIREYRL TGRARRLAIRPGSPMIGQRLDG LULRERYGNANIGVER WRFFR RVYNVNGVSEFRARDVSLIAM SAADVDLRQFCSERLLEPDGIA RSADIPESELIGESVREIGERTRY GLNYVNGVSEFRARDVSLIAM SAADVDLRQFCSERLLEPDGIA RSADIPESELIGESVREIGERTRY GLNYVGLTRTGVARADSA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24768	55136	A	24906	834	1993	EEIEVARI.RDQKSAEGGIQNRE YWPGRSGHR*SPHAPIIGTQFLQ ALFQTWDQFVFALSACGRRHP TDGRRQNPAVSGHSVAARQPA HSQTDEASG
24769	55137	Α	24907	3519	5161	
24770	55138	A	24908	374	1891	KWKLKRPDSNIDHRRVPNLET WFSRHDKTRPTSKNPSDYQAG DIVSWRLDNGLAHIGVVSDGFA
						RDGTPLVIHNIGAGAQEEDVLF NWRMRQCQHNQHNTATFNGP RILAEOPPGKEYAANGNGOLYP
						MSSLNIKQGSDAHFPDYPLASP SNNEIDLLNLISVLWRAKKTVM
						AVVFAFACAGLLISFILPQKWTS AAVVTPPEPVQWQELEKSFTKL RVLDLDIKIDRTEAFNLFIKKFO
						SVSLLEEYLRSSPYVMDQLKEA KIDELDLHRAIVALSEKMKAVD
						DNASKKKDEPSLYTSWTLSFTA PTSEEAQTVLSGYIDYISTLVVK ESLENVRNKLEIKTQFEKEKLA
						QDRIKTKNQLDANIQRI.NYSLD IANAAGIKKPVYSNGQAVKDD
				· ·		PDFSISLGADGIERKLEIEKAVT DVAELNGELRNRQYLV\EPITK AHVNDVNFTPFKYQLSPSLPVK
						KDGPGKAIIVILSALIGGMVAC GGVLLRYAMASRKQDAMMAD
24771	55139	A	24909	1111	1895	PLVNFSNSSHPVSNPSIQLSSITL PLVENESRARA\VKDQMDAIDH QTGVRPLVFYSIVLPEIRAIILQS
						EGFCQDIVQALVAPLQQEMKL DPTPIAHRTHGLNPNNLNKYDA
		ļ				RIAAIDYTLAHDDGISLRNLDQ AQVILLGVSRCGKTPTSLYLAM
						QFGIRAANYPFIADDMDNLVLP ASLKPLQHKLFGLTIDPERLAAI REERRENSRYASLRQCRMEVA
						EVEALYRKNQIPWINSTNYSVE EIATKILDIMGLSRRMY
24772	55140	Α	24910	77	434	EKKRGREDKPGTMATFPPATSA PQQPPGPEDEDSSLDESDLYSL AHSYLGPLIMPMPTSPLTPALV
						TGGGGRKGRTKREAAANTNRP SPGGHR/ERKLVTKLQNSERKK RGARRLRQSWR
24773	55141	A	24911	31	295	

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, :=possible nucleotide insertion)
				sequence		
24774	55142	A	24912	75	426	EKKRGREDKPGTMATFPPATSA
24,,,4	55142	ľ.	24712	1,3	420	PO\OPPGPGGRDSQPWMDLDLL
						/YSLAHSYLG\GGGRKGHTKRE
	1	1				TAANTOPPOPLAGHERKTVDP
						KLONSERKKRGARALROELEM
						RRRPWTLTPSO
24775	55143	A	24913	1	1107	and Highley
24776	55144	Α	24914	1	1468	
24777	55145	Α	24915	I	2463	
24778	55146	Α	24916	1	603	
24779	55147	Α	24917	118	1087	IRSKKMTTQTVSGRRYFTKAW
						LMEOKSLIALLVLIAIVSTLSPN
		l				FFTINNLFNILQQTSVNAIMAVG
1		1				MTLVILTSGIDLS/VAASIVGIEV
						NALVAVAAALALGAAIGAVTG
						VIVAKGRVQAFIATLVMMLLL
						RGVTMVYTNGSPVNTGFTENA
						DLFGWFGIGRPLGVPTPVWIMG
l						IVFLAAWYMLHHTRLGRYIYA
						LGGNEAATRFSGINVNKIKIIVY
						SLCGQLASLAGIIKVARLSSAQ
						HTAGTGYELDAIAAVVLADDR
				ĺ		OSGGDFAGGAGROOKAVITTT
İ						GHLEYEHEKTGYPGFRCCAKR
		1				HRQCECDGKRHHRAGGLHA
24780	55148	A	24918	1	1286	MHEREPGLRAALAGOLEFRVG
						LGLGVWTLRLMELLRORLELV
		1				VYTASDGQEYEMVSRLNWVN
						YLVLEDVHIILPDQMRGITGCY
						ATHPHERRLHAGTVHRONIVN
İ						NPPDCVPRDAGEAEVAPDPSLA
						SLQLQTPELPGLISCHNSIQVSN
						EARLFELLICOWRDFRKRVIAA
		1				KSMQGRELEGGQRGQDKATNP
						FRVLALKGSGDPDPHGAGRQG
						SDLLLHPVSDAQVLQHFFQRRA
	l			1		LLSSPWDYHDVVGLCAQEPCG
1	1	1	1	1	1	YIMELFIIKHQCSLAQTAGSLAP
		l		1		EQKKELEMTVRYYISSADLTAE
l		1				KFATAIRN/HWHVENKLHWRL
l	1					DV*RQFDKHCSTADSTRILASF
l		1				VRHVNAALDVRVVFASGSETT
l						HGDGARRYLRCVDVDWRRRT
	1					AVAQADKSARACHFHYAGYH
					1	HHEHFADPVSVGAKYHTVFST
				L	L	PHILLIADI VOVOAKIIII VESI

## WO 01/075067

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	1	
24781	55149	Α	24919	417	861	AKGPCPDRNHSSDDKDVIAIDG
24761	33143	n	24515	1117	1001	KTLRHSYDKSRR\RGAIHVISAF
	1					STMHSLVIGQIKTDEKSNEITAI
		İ				PELLNMLDIKGKIITTDAMGCO
l	1	l	ŀ			KDIAEKIOKOGVIFSLCPASIIDD
						KKVTDLAVAVQGTLKTGSGGI
						LPPGGALKKISEKL
24782	55150	A	24920	3	1767	HISIIPDYR*AWKVEHKLSGILL
24/62	133130	^	24920	ľ	11707	LTIFAVISGAESWKDIEDFGKTH
						LDFWKOYGDFENGIPVHDTIAR
ŀ	l		İ			VVSCISPAKFHE\CFINWMRDCH
ŀ						SSNDKDVIAIDGKTLRHSYDKS
					į.	RRRGAIHVISAFSTMHSLVIGOI
						KTDKKSNEITAIPELLNMLDIKG
	ŀ				ŀ	KIITTDAMGCQKDIAEKIQKQG
					ľ	GDYLFAVKORNGAVENRAKTO
į .						TRIAFGDHFFRROVSORIRLFCH
1						RLATOFAARLRMFIATGOPMN
	1		1			GLRLLLFIVNKOAALLVNKLDA
						QMAAFQQLPGRGRQATVRRE
						MDLCAFAERMGLPVSRPGDIOE
	1	i				FHAGCRTKFDAGAFRDKHHVA
Ì	1				ł	CFGDHGLAFVANMOLPFOHHV
İ						DFVVVQGPREECVLPLHKTGG
	1					HAAAVVSGHKISQAHAMRLCA
ŀ			i			VETTKLDATGINHFDAAFAGHR
ŀ	1	l				FESSVDCACSATVSFITFPSVCD
ł		l			l	RFGNWLRSGRCSHAARKNEITL
	1	l				TNTRRSLRPKRNTERHDMQTR
		1			l	ATVIAAGEKLLPYAETLMSTW
}	i		İ			OAARKEVAHTSRHNEFSIGASA
1	1		1			SLWDMVMTSAGAVVTTVVVA
					İ	VRPSSFYILCSHKKKRTLHTKR
			1		ĺ	EKTVDRRKEYETRAKRSETDT

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence	1	
24783	55151	A	24921	110	1282	SNTPYLKENKMELKKLMGHISI
24763	122121	^	24921	1110	1202	IPDYRQAWKMEHKLSDILLLTI
		1				CAVISGAEGWEDIODFGETHLD
		l l				FLKQYGDFENGIPVHDTIARVV
		l		l	İ	SCISPAKFHECFINWMRDCHSS
		l				DDKDVIAIDGKTLRHSYDKST/
		1			l	RRRGAIH\VNNAFSTMHSLVIG
		l			ł	OIKTDKKSNEITAIPELLNMLDI
		1				,
						KGKIITTDAMGCQKDIAEKIQK OGGDYLFAVKGNOGRLNKAFE
		l		ĺ	}	
ł		l				EKFPLKELNNPEHDSYAMSEKS
		l				HGREEIRLHIVCDVPDELIDFTF
		1				EWKGLKKLCVAVSFRSHAEQK
						KEPEMTVRYYISSA\DLTAEKFA
		1				TAIRNHWHVENKLHWRLDVV
		l				MNEDDCKIRRGNAAELFSGIRH
		1				IAINILTNDKVFKAGLRRKMRK
		_				AAMDRNYLASVLAGSGLS
24784	55152	В	24922	258	507	
24785	55153	Α	24923	1	3060	MTEIEENMLLPISGQELPIRWLA
		l			l	QHGSEKPVTHVSRDGLQALHIA
		l			i	RAEELPAVTALAVSHKTSLLDP
		1				LEIRELHKLVRDTDKVFPNPGN
		l				SNLGLITAFFEAYLNADYTDRG
		l				LLTKEWMKGNRVSHITRTASG
l		1				ANACGGNLTDRGDSFVHDLTS
		1			l	MARYVATGVLARSMGLVIYSFI
		1				QHTLTLEEYREITAFLFREFTHA
		1				AAARYTGEQETLPLSPQWILRQ
		l			ŀ	CKEVASLCDGDTFSGEQLNLM
						LQQREWREGFLAER
24786	55154	В	24924	56	967	
24787	55155	В	24925	1	1776	
24788	55156	A	24926	1	864	
24789	55157	Α	24927	424	513	
24790	55158	IA.	24928	1	5049	l i
24791 24792	55159	В	24929 24930	1	939 1837	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24793	55161	A	24931	I	2249	MLPDIARFLPRLANDQIEEFTW YVNHFTYSNAIKIFCYQMFLAS TQARASGAPLQRLFRAIGGNA WERPFRMVFCSAAFSEESSPFR
						ETNYEPQLFLGFATDYRFAGW TLRDVEMGYNHDSNGRSDPTS RSWNRLYTRLMAENGNWLVE VKPWYVVGNTDDNPDITKYM GYYQLKIGYHLGDAVLSAKGQ YNWNTGYGGAELGLSYPITKH
						YNWNTGYGGAELGLSYPITKH VRLYTQVYSGYGESLIDYNFNQ TRVGVGAIGVNVAQAEVLNLE SGAKQVLQETFGYQQFRPGQE EIIDTVLSGRDCLVVMPTGGGK SLCYOIPALLLNGLTVVVSPLNS
						LKEKSANPFEFTFAAFAHAFHW VIQAIGMINTPADGSTHKQART SWLPYSSLPVLSVSTQDEKAKK KMRFMQRSKDSLAKWLSAILP
						VVIVGLVGLFAVTVIRDYGRAS EADRQALLEKGNVLIRALESGS RVGMGMRMHHVQQQALLEEM AGQPGVLWFAVTDAQGIIILHS DPDKVGRALYSPDEMQKLKPE
						ENSRWRLLGKTETTPALEVYRL FQPMSAPWRHGMHNMPRCNG KAVPQVDAQQAIFIAVDASDLV ATQSGEKRNTLIILYALATVLL
						ASVLSFFWYRRYLRSRQLLQDE MKRKEKLVALGHLAA/GVAHE IRNPLSSIKGLAKYFAERAPAGG EAHQLAQHVQIRRFHAHHPAS

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24794	55162	A	24932	261	2115	TLVFCWHAKLLSHGVFMDIISV
						ALKRHSTKAFDASKKLTPEQAE
						QIKTLLQYSPSSTNSQPWHFIVA
					1	STEEGKARVAKSAAGNYVFNE
			l			RKMLDASHVVVFCAKTAMDD
					i	VWLKLVVDQEDADGRFATPEA
			ŀ			KAANDKGRKFFADMHRKDLH
					l	DDAEWMAKQVYLNVGNFLLG
			i			VAALGLDAVPIEGFDAAILDAE
	İ					FGLKEKGYTSLVVVPVEDLAGI
		1	İ			EIDHTTSMVMIFGIIFLTAVVVH
						IILHWVVLRTFEKRAIASSRLWL
						QIITQNKLFHRLAFTLQGIIVNIQ
					i	AVFWLQKGTEAADILTTCAQL
						WIMMYALLSVFSLLDVILILGQ\
						KFPA\AFQLPLKG\IFQGIKLMGP
						ILVGILMISLLIGQSPAILISGLGA
						MAAVLMLVFKDPILGL\VAGIQ
1						LSANDMLKLGDWLEMPKYGA
				ŀ		DGAVIDIGLPTVKVRNWDNTIT
						TIPTWSLVSDSFKNWSGMSASG
						GRRIKRSISIDVTSIRFLDEDEM
					1	QRLNKAHLLKPYLTSRHQEINE
						WNRQQGSRESVI\NLRRMP\NIG
						\TFCAYLNK\YLGTH\PRIRKDM
						TLMVRQLAPGDNGLPFEIYAFT
1	ľ				1	NTVVWLEYESIQADIFDHIFAIV
i						EEFGLRLHQSPTGNDIRSLAGA
						FKQ
24795	55163	Α	24933	1	757	
24796	55164	A	24934	8	343	
24797	55165	A	24935	I423	1761	A CONTROL OF THE CONT
24798	55166	Α	24936	1	851	MKGDTKVINYLNKLLGNELVA
			ŀ			INQYFLHARMFKNWGLKRLND
			İ			VEYHESIDEMKHADRYIERILFL
			i			EGLPNLQDLGKLNIGEDVEEML
		ļ	ŀ			RSDLALELDGAKNLREAIGYAD
						SVHDYVSRDMMIEILRDEEGHI
						DWLETELDLIQKMGLQNYLQA
	l		l			QIREEGKSIYQEQTTAMKRPDY
		1				RTLQALDAVIRERGF/EARGTK
1		l	l	l		AVHYTISRLTAH*ATGKYVRAA
1			l	l		AVGAYRTAAPDGTRAKTAGTA
1		1	l	l		APGGVAGRRVAGR*TNRFDSA
						AAFTGGQRRQSGDVVASCTGS
	L		L	L	l	CVG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
2.4700	100160	<u> </u>	0.1000	11645	2009	I PVFIOFVEGMCGTANEPACLLV
24799	55167	Α	24937	1645	2009	
						NQSPAIYFPGKIAAIWQNPRHV
						ESGSM*TL/IDSD*VSPTECLSPR
		l				HHYMPFEEAARGYEIFEKREEE
1						CRKVILVPGAQSAEAAQKAVS
						GLVNAMPGGTI
24800	55168	A	24938	1	3936	
24801	55169	Α	24939	1	2240	MRSPMTRNSRNGKHDYKPPPR
		1				KRRQPPHDHTAPNSTPARNQRO
		1				RPHRPNRQAGQREQRANNPTP
			1			NTPSPPPTNDPAEPHQTRHDSR
						QPHNTPPAYAELVAAALLTTDT
						NVTRRRLRYMQPDTKLCHSAG
	1					KSIYQEQTTAMKRPDYRTLQAL
						DAVIRERGFERAAQKLCITQSA
		1				VSQRIKQLENMFGQPLLVEDET
1		1				RTQERLRRGEVVGAVSIQHQAL
		l				PSCLVDKLGALDYLFVSSKPFA
1		l		ļ		EKYFPNGVTRSALLKAPVVAFD
		l				HLDDMHQAFLQQNFDLPPGSV
		l				PCHIVNSSEAFVQLARQGTTCC
1		l				MIPHLQIEKELASGELIDLTPGL
l	i	1		İ		FORRMLYWHRFAPESRMMRK
		1				VTDALLDYGHKADDIILRITAT
		1				AICGSDLHLYRGKIPOVKHGDI
						FGHEFMGEVVETGKDVKNLQK
1						GDRVVIPFVIACGDCFFCRLQQ
1	1	1				YAACENTNAGKGAALNKKOIP
						APAALFGYSHLYGGVPGGQAE
		1				YVRVPKGNVGPFKVPPLLSDD
1		l				KALFLSDILPTAWQAAKNAQIQ
	1	1				QGSSVAVYGAGPVGLLTIACAR
	1					LLGAEOIFVVDHHPYRLHFAAD
1		1				RYGAIPINFDEDSDPAQSIIEQTA
1			ł			GHRGVDAVID\AVGFEAKGSTT
1		1				ETVLTNLKLEGSSGKALRQCIA
1		1				AVKRCGIVSVPGVYAGFIHGFL
1		1				
1	ŀ	l				FGDAFDKGLSFKMGQTHVHA
		<u>_</u>				WLGELLPLIEKGLLKPEEIVTHY
24802	55170	Α	24940	687	792	TPTLTRPLRHPVAYRATLK*NR
				L		PAKVSQLCCTNA
24803	55171	Α	24941	3	510	AGDMVWLCVRTQISY*IPRTHP
		1			1	AVGIPRERPLRGADTPQRGASH
1		1			1	RIRGEGKRK*KRWALLCCPGCN
1	1	1				I*RHFCHHN*ERYFW\IYWVEST
1	1	1			1	GAVKHLQNTGHWLNAGALPV
1	1	1			1	RRPRKSVTGSIARQ\VSVPISAR
1				1		ALLACPCGQCAGYATQFVMKS
		1				ITORFNVACRTRMYAVG
24804	55172	l <sub>A</sub>	24942	189	563	
£4004	100172	l''	2772		L	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24805	55173	Α	24943	1224	1772	GSPSVSLATSSCSIKVQNPGISR
						VVAGSSAIISNVSPGCILRTVLR
		ŀ				TIITGSGQRRPSASSVWSGHTSC
		l				VFRTANFSSGMVNLSVSVTNLS
	Į.					PSIVPRIKFIAGDPINPETNIFAG
1	i .			l		WL/VNGFRRTYLLNNPVFHNH
l	I	1		i		NPRRHGHRFGLIVGHVDHGGV
				l		QFLMQLADLCAHLHAQLGIKV
24806	55174	Α	24944	1541	3675	LEHKRKISMRRNWMHSHTRY
	i i					WRFLDFPTLRGGRWMR*SGEK
	1	ı				LNPISTPLRANCLALAQRATRL
		1				APMSFTWGMPTSQPDCAVPGR
		1				RALPPELLCLQRSSQSRFAGTAI
			1	1		T
24807	55175	В	24945	1	1497	
24808	55176	Α	24946	650	739	
24809	55177	Α	24947	4447	4804	RQNDRYSGRERHEITPLCW*IPR
						IAWALWLRSLRRGPQEKSLHN
						RVARTISWPYQARHPCRTLLGG
		1				HGFFLHIVLSQLPCLSYPLSSFA
						GLLARSSMFLKLAARFLCCRRS
						YAASPLSI
24810	55178	Α	24948	1	3774	
24811	55179	Α	24949	431	949	NYAIYHQNHGVNISPGLGKTHN
l	1					YRQQHCGITYYWRDLMPANK
				1		NYKQRWITPGSASFYRSSHKAL
				1		IRLLATRLPAFPWGRRSAWRWP
		1				VRY*IRVPSSGSYHLEISMRWVI
		1			1	FPQPKALTPSNLHWKTA*LWS*
					1	KTLSFVCWIATVLPARFWKLTD
						PWLAILPCANIYWQASAPD
24812	55180	Α	24950	337	648	

SEQ 1D	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
		<u> </u>				
24813	55181	A	24951	1	1430	MFTYPHSKFRVNRFVGQVLLIE
					i	FSILLDGGRTGLGHSLSVWCISG
İ			ĺ			DCSSFLQGKPTSLPLRKPSAIICS
ł	ŀ					DSLSGSADLLRNHNFVCTAADT
İ						FTTDITPEAGDARLIGIAQTVRH
						LEDPMRAWKSIHKSCEILDPIHR
l					l	APVKENTHPARKINISDPAAGA
						VARIRYEGVLSVCRASASAYVS
l	l	1	ŀ			RSRTITVACWYVSNTGVTGATI
	1	l				GVKRQHRHARLFSGDGAPGFG
İ						TGKRDIRQLRRLVRIQIQRTFFA
						AVLHVPTPWSRAIDAAALLTVI
		l				DVKVLTACACQACRTLALRVA
	ŀ				1	QVIDRCSNPGDIMSSVARRYWF
l						ISRWSAKRDCAVSYDQDPGQA
						ORSCRSCLVAVDRGLWLSAYC
i .						RSLMTQTLLPDDVPRILPNASSP
i						SRLRCL\HIWVITACWICANSSG
		1				FIARNL/GLCGWEKRLVKDLTRI
1		1				ODTGRAKEILGATATLEFR\R*T
1		l l				PTLTRPLRHPVAYRATLK*NRP
1		ı				AKVSOLCCTNA
24814	55182	В	24952	1	1295	ARVSQUEETIVA
24815	55183	A	24953	3151	3207	LQWLID*WCL**TFDLRFA
24816	55184	A	24954	2740	3917	NGRSVWAVKLELFRVKPLDIPO
24810	33164	l <sup>A</sup>	24934	2740	3917	QRVGQQHVLLKAFNRRLAMQF
		l				
		l				IQLIATDVTEILPLHVVEDGTDS
		l				LYCLALSHLRCGRLSLRWRREL
1		ı				PRGLGVNVSWCYPFKAQGQCA
1		1				AEITDKAGQREIFGFARVKENA
1		1				GYVIYECYQPDDGDKLIRELPF
1		1				SSLIFARQWFVVGELLQHLPPE
1		1				DRITPIVGMLQGVVEKGGELRV
l		1				EVADTNESKELLKFCRKFTVPL
1		1				RAALRDAGVLANYETPKRPVV
		1				HVFFIAPGCCYTGYSYSNNNSP
		1				FYMGIPRLKFPADAPSRSTLKLE
1	1	1	1			EAFHVFIPADEWDERLANGMW
						AVDLGACPGGWTFQLVKRNM
		1				WVYSVDHGSMVQ/SLWITTGT
						CWGTRFHSARGOKFPEVLOGS
						QTLPAATDPPGMQWLASGVLE
24817	55185	A	24955	1	519	
24818	55186	A	24956	510	731	
27010	55.00	<u></u>	1-1,550	1	1	l

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24819	55187	Α	24957	ī	1055	MRLCPEKTTKAKRVKAVLAWS
		1				KEEPWLSEWSEEVLEMGYKKL
ĺ	ŀ					GDFSTAGAILETEIRPSPASAFTL
		1				DSQPPESNNLSEAKYAPNNIESI
		1				AKYFAPKSPMFSESPSGTALDS
1	l					FNRNERRRWPLWPNKSGFGSD
		1			1	LTLSKIGISSPDIALERLYEQAR
		1				KLSGKFRALSELDESARSGVLK
ĺ	1	1			}	VLCAFAYQDYSRSAASTRKCD
		1	l			CCDGGGFTEAQVFTNKVSYPW
ŀ	l	1				GKPPYWSKMSRAVRPSDWESW
			l			TQAREVVRVKCKPCNGKGVIS
						NSCRCHGKGKVLAKHESDRRG
		1	Ì			VPVMKACDRCGGRGYARLKFS
		1	İ			SV\MRALILLRRLKKTAAYEQL
	1	l				QPLFEELVAECHKQESMADSIL
						SKVTR
24820	55188	Α	24958	361	725	
24821	55189	Α	24959	1100	1738	WQRSTLPLLFESAMQTGVNAG
		1				NPVAW*KSAGYGHIKLYDAKN
	1	l				RL\LVQSTALWASQLVQYLSTF
	ł	1	ŀ			FRKNLKRPSEFVTLADEIEHVN
						AYLQIEKARFQSRLQVNIAIPQE
	İ					LSQQQLPAFTLQPIVENAIKHGT
						SQLLDTGRVAISARREGQHLML
	ł					EIEDNAGLYQPVTNASGLGMN
					i	LVDKRLRERFGDDYGISVACEP
		Ļ.				DSYTRITLRLPWRDEA
24822	55190	A	24960	1	2043	WI WALL BUDGETOUR DE LE SOOT
24823	55191	Α	24961	2164	2487	KLKRIARHNFHGIRDLHLFQQV
1		1	İ	1		NRFSSGFPTTYWLMMADHVHH
		l		1		LKTYSTCRIEGGHRFLKDHRHS
		l		1		CAAPLTKLFFPRVKNIFSQQGN
	L	Щ.	L	L	l	LPPMYAITARR*QTHNQHCGD

SEQ ID		Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24824	55192	A	24962	128	2574	OVVVVMAMVAKVVIDIVVVV
24024	33172	<u> </u>	24702	120	2574	GVEMIRVKVVAIVVEVVVVEV
						VVVMVAWWW****WW*/DMG
						EARWWWEYGVWEVEPEIILNA
						POOYLLAGIGDTLAKWYEAVV
						LAPOPETLPLTVRLGINNAOAIR
1						DVLLNSSEQALSDQQNQQLTQS
i						FCDVVDAIIAGGGMVGGLGDR
						FTRVAAAHAVHNGLTVLPOTE
						KFLHGTKVAYGILVQSALLGQ
						DDVLAQLTGAYQRFHLPTTLA
				i		ELEVDINNQAEIDKVIAHTLRPV
						ESIHYLPVTLTPDTLRAAFKKYS
						OSCFVIRLGGDFIHOLAENHIVI
						FVNNHNGTRCOTFOWAISNCH
						TVSLQKFSVRKFLSDLFAIGGR
				ľ		RGVRPPTEAPGHOKSDNOAHV
						SVSFDGLDLVFFVVNOWAGVV
						FCFDLFSRKRGRFSVWRNKGFG
						RYCQRLANFRDRQAFQLLNQIH
						FTRCKQVDNGGTEFKATHFLAF
						QQCHRTVVFPGRPAIKTRVERL
						DRYRNRMEKSNGQANLRVHG
						GTWEKLVLQLDSAGMVNRVSS
						VGAPLREPIPTEGKGAKEDTWG
				l		NLSSNDPLLEAFGGQFGIAISMH
				1		QTLATRGGKRNARPTTRPRFHR
						GTNPRGIAPHIGTANARYSTGH
		-8				CPVALPFGLRGPPLARENVATA
				l		TKGHESASRETEVVAVAKLTSR
						VLLICDATHALPIQVLRNDIRVI
						AAGNGCTDDLQLEHLPANTDIR
24825	55193	Α	24963	117	335	PGLAPATRRSTPGRFNADAGAD
						*RRAAGTGAGRAKRRGVGIHR
				l		CLADPFLADCRNAGLRGAALC
						GTAAASGK

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon fur last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \*=possible nucleotide insertion)
24826	55194	A	24964	208	1903	YLAIKTERCLHDEFTDEQLSRA VWIYOKRAIAAA QTKLPPAFGL PGAKHILFROHCSESDVQLAA ESGDDRSVVIGVGGGALLDTA KALARRLGIPVA VPITAATCA AWTPLSVWYNDAGOAL HYEIF DDANFMVLVDPEIILNA PQQYL LAGIGDTLAKWYEAVVLAPQP ETLPLTVRLGINNAQAIRDVLL NSSEQALSDGONQQLTOSFCDV GDAIIAGGGMYWGLGDRFTRV EAAHAGPNGLTGLPOTEKFLH GTKVAYGILVQSALLGQDDVL AQLTGAYQRFHLPTTLAELEVD INNQAEIDKVIAHTLRPVSEDIDVSLAA RQLAQALAEGNGVAWESIDA WQIPSSLIAEMPASVALHYAVL PLRLENDELIVOSEGDIDVSLA ALTRKVGRKVRYVIVLRGQIVT GLRHIWYARRGHDPRAMLYN AVOHQWLTEQAGEIWRQYVP HQELFAEILTTLGHINRSAINVL LIKHRESSLPLGKFLVTEGVISG ETLDRVLTIGRELQVSMOSLLI
			21011		205	KAGLNTEQVAQLESENEGE
24827 24828	55195 55196	В	24965 24966	69	205	
24828	55196	В	24966	1	721	
24829	55197	A	24967	1	1703	
24831	55199	A	24969	2853	2889	LIIGLDLPAVSAIFQRALRFTHV GIADGGAYVVQR/HRPDEPRRR CCSE
24832	55200	A_	24970	297	473	
24833	55201	A	24971	1179	2148	CQVLRPRTKRLNVLFLHRNOM QWRORELVINMCFVIACAKR RAPFVQAAEVRPOQHDDEFP LAIWQTGSGTQSNMNMSVLA RASELLGGVRGMERKYHPND DVNKSQSSNDVFPTAMHVAAL LAKQLIPQLKTLTQTLNEKS RAFADIVKIGRTHLQDATPLTL GQEISGWPASRSGTGSWRYSGG WYTKYPSGVCASRSKMNWYST RASDGGHASDA VRTAQAADRE LRAPVRDELNASITSKTRLAEST SLAEWYKASPHDDDKPILLPGE WEVNTRRERQKQGIPLDAGSW QAICDAARQIGMPEETLQAFCQ Q

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24834	55202	A	24972	166	365	NPFSNSPWETYPVAFTIPLALFM GIYLRYLRPGRIGEVSVIGLV/SP DFRHYLWRLGGRKSDLGTVL
24835	55203	А	24973	83	362	KPPCFLQASVTCWLRLPFASMII LPPCACS*ST*ESIRPAVVGPREP EGITFRFGRTRVVNRMIFDVL RHLFTVIHPLFQFGVTYITTDDN
24836	55204	A	24974	584	853	ROLL TVIIII EL QUOVITTITIBEN
24837	55205	А	24975	1	3017	MSQDPFQEREAEKYANPIPSRE FILEHLTKREKPASRDELAVEL HIEGEGQLEGYAPVLAPESGILI MRPTLTMPALTKFVDGTGPVW TGNLPFPLFITIACGVSGFIBALI SSGTTPKMLANEGQACFIGYG MLMESFVAIMALVSACIIDPGV YFAMNSPMAVLAPAGTADVV ASSAQVVSSWGFSITPDTLNQI ASEVGEQSIISRAGGAPTLAVG MAYILHGALGGMMDVAFWYH FAILFEALFILTAVD
24838	55206	A	24976	3	206	EPDFLPL**RRNHEQNYRDGKC TGSYRSLRTGR*SGQYDHHLRS DPGKSENGRSTGRRRCTGTSVA G
24839	55207	A	24977	3	982	GDAEPPVFLPRJKLDHEVSLDH YQHRFYLRSNRHGKNFGLVRT RMRDEQQWEELIPPRENIMLEG FTLFTDWLVVEERQRGLTSLRQ INRKTREVIGIAFDDPAYVTWIA YNPEPETARLRYGYSSMTTPDT LFELDMDTGERRVLKQTEVPGF YAANYRSEHLWIVARDGVEVP VSLVYHRKHFRKGHNPLLVYG GGYGYAIDADF\$FSRLSLLDR GFVYAIVHVRGGGCKINLVVD NGKIVRAEAAGGKTNQGTLCL KGYYGWDFINDTQILTPRLKTP MIRRORGGKLEPVSWDEALNY VAERLSAIKEKYOPDAIQTTGSS

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	or pepude sequence	deletion, possible ancientide insertion)
24840	55208	Α	24978	1339	3640	SPLSTAGISICGPLLIITDVDLGT
						TVAGIKGMIATGEGTTEWNSV
						MVAMLLTLIPPVVIVLVMQRA
		1				VRGLVDMRNKMAGLKLOAVT
						KSWDGKTQVIKPLTLDVADGE
						FIVMVGPSGCGKSTLLRMVAG
		1				LERVTEGDIWINEQRVSELRLIC
	1					YMSRKMGGGKNRREPHIDVKF
	i .	ļ				GIFGLSITPEFSEPLAEDFLSRRQ
		1				YGIRFRNLETGNWYPELLDNVI
	1		l	1		PSFVWANDSWIFYYVRKHPVT
	ł	1	ŀ	l		LLPYQVWRHAIGTPASQDKLIY
	1			i		EEKDDTYYVSLHKTTSKHYVV
	l					HLASATTSEVRLLDAEMADAE
						PFVFLPRRKDHEYSLDHYQHRI
	1					YLRSNRHGKNFGLYRTRMRDE
						QQWEELIPPRENIMLEGFTLFTI
			Ì			WLVVEERQRGLTSLRQINRKTI
						EVIGIAFDDPAYVTWIAYNPEPI
						TARLRYGYSSMTTPDTLFDLDI
						DTGERRVFKPPE/V/PGFYAA/SY
						RREPRW/IVSADGVKYSFV\APR
	İ					QHFRRDAPVLSFLLFSDRAPFR
						NNGPRVVLLLGYYHPELLGVT
	ŀ	i	İ			YSQLSVHEARPKPLPIPHSRHRI
						VPTRNEMAQEKERPAGSQSLFF
						GLMLIEILSNYPNGCPLAHLSEI
	1					AGLNKSTVHRLLQGLQSCGYV
			l			TTAPAAGSYRLTTKFIAVGQKA
						LSSLNIIHIAAPHLEALNIATGET
						INFSSREDDHAILIYKLEPTTGM
						LRTRAYIGQHMPLYCSAMGKI
24841	55209	A	24979	1	248	
24842	55210	Α	24980	839	1625	PDAFILTTKPAGAIPTSTSITRPI
			i			FCPSFAPCENATPAAET/HQRN1
						RPEWRFTFAFFLFAIFRNAVNT
						VTMSIRKGIGVHHNAANLVPW
				1	1	MTKFFQRFGIAFQSRGAVFHQF
						HHAGYAGQRGKTALAEQRLVI
						GFGINKQMRAVPLIGEVYQFSG
						*CQRQSFTADGFTDSNAFNNIS
						GNSTARDKLFIAGYGNRYFHC
						GVNAQIIFMQELGDFRHFVIF/A
						QKGNEYHGDTRISPGRFRRGHF
						PLGALRLAASVERSGNGHRA
	55211	Α	24981	2538	2864	GSSLINSKVMP/SF*QQNRKFAY
24843	33211				1	
24843	33211				1	KRAGVATDEFHAESFVRGRAP
24843	55211	ļ .				KRAGVATDEFHAESFVRGRAP MKRKL*LFMQLGGDVIQPFDGI
24843	55211					
24843	33211					MKRKL*LFMQLGGDVIQPFDG